

A Dissertation

entitled

An Investigation Into the Fate of a C5'-Uridinyl Radical

by

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Submitted to the Graduate Faculty as partial fulfillment of the requirements for the

Doctor of Philosophy Degree in

Medicinal Chemistry

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Constituents of the endogenous exposome include reactive oxygen species, which cause oxidative damage to nucleic acids, proteins, and lipids. It has been shown that such damage to RNA plays a role in the initiation of neurodegenerative diseases, such as Alzheimer's. To investigate this phenomenon at the molecular level, the synthesis of a modified uridine was improved upon and subjected to photolytic activation, generating a C5'-uridinyl radical mimicking the hydrogen atom abstraction normally observed under conditions of oxidative stress. This was accomplished using a *tert*-butyl ketone at the C5' position, which undergoes Norrish Type I cleavage. Studies were performed under anaerobic conditions in the presence or absence of reductant at varying pH to emulate the diversity of the cellular environment, and it was found that the base release product uracil was predominant. Aerobic conditions were also utilized, which led to the formation of a C5' aldehyde. The C5'-uridinyl radical precursor was further derivatized for its incorporation into oligoribonucleotides. Development of a new automated RNA synthesis method in the 5' to 3' direction using 5' H-phosphonates was initiated. This included the synthesis of the building blocks required for the method development.

“God, grant me the serenity to accept the things I cannot change,
courage to change the things I can,
and wisdom to know the difference.”

- Reinhold Niebuhr

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List of Abbreviations

^1H NMR	Proton NMR
^{13}C NMR	Carbon NMR
^{31}P NMR	Phosphorus NMR
A	Adenosine
Ac	Acetate
ACN	Acetonitrile
ATP	Adenosine Triphosphate
Bz	Benzoyl
BzCl	Benzoyl Chloride
C	Cytidine
C2	Carbon 2
C4	Carbon 4
C5	Carbon 5
C6	Carbon 6
CPG	Controlled Pore Glass
d	Doublet
dd	Doublet of Doublets
DCM	Dichloromethane
DMAP	4-Dimethylaminopyridine
DMSO	Dimethyl Sulfoxide
DMTrCl	Dimethoxytrityl Chloride
DNA	Deoxyribonucleic Acids
EA	Ethyl Acetate
ESI	Electrospray Ionization
EtOH	Ethanol
FaPy	4,6-diamino-5-formamidopyrimidine
FaPyA	FaPyAdenosine
G	Guanosine
GSH	Reduced Glutathione
GSSG	Oxidized Glutathione
GUA	Guanosine, Uridine, Adenosine
HPLC	High-Performance Liquid Chromatography
HRMS	High Resolution Mass Spectrometry
Ibu	Isobutyryl
IR	Infrared
LCMS	High-Performance Liquid Chromatography-Mass Spectrometry
m	Multiplet
MeOH	Methanol

mRNA	Messenger Ribonucleic Acids
MS	Mass Spectrometer
NMR	Nuclear Magnetic Resonance
nt	Nucleotide
P (III)	Trivalent Phosphorus
P (V)	Pentavalent Phosphorus
Pyr	Pyridine
O	Oxygen
OPC	Oligonucleotide Purification Cartridge
q	Quadruplet
Q-TOF	Quadrupole Time-of-Flight
RNA	Ribonucleic Acids
RP-HPLC	Reverse Phase-HPLC
rRNA	Ribosomal Ribonucleic Acids
s	Singlet
SAM	S-Adenosyl Methionine
sep	Septet
siRNA	Small Interfering RNA
t	Triplet
<i>t</i> -BuLi	<i>Tert</i> -Butyllithium
TBAF	Tetrabutylammonium Fluoride
TBATB	Tetrabutylammonium Tribromide
TBDMSCl	<i>Tert</i> -Butyldimethylsilyl Chloride
TBDMSCN	<i>Tert</i> -Butyldimethylsilyl Cyanide
TEA	Triethylamine
TEA·3HF	Triethylamine Trihydrofluoride
TEAA	Triethylammonium Acetate
TEAB	Triethylammonium Bicarbonate Buffer
TFA	Trifluoroacetic Acid
THF	Tetrahydrofuran
TLC	Thin Layer Chromatography
TMSCl	Trimethylsilyl Chloride
tRNA	Transfer Ribonucleic Acids
U	Uridine
UV	Ultraviolet Light

List of Symbols

\geq	Greater than or equal to
'	Prime
$^{\circ}\text{C}$	Degrees Celsius
α	Alpha
β	Beta
δ	Chemical Shift
\AA	Angstrom
μL	Microliter
μm	Micrometer
cm	Centimeter
eq	Equivalent
hr	Hour
h ν	Energy
J	Coupling Constant
L	Liter
M	Molar
mAU	Milli Absorption Unit
min	Minute
mL	Milliliter
mM	Millimolar
mmol	Millimole
mv	Millivolt
Mz	Magnetic Field Strength
m/z	Mass per Charge
N	Normal
nm	Nanometer
nmol	Nanomole
W	Watt
Ac_2O	Acetic Anhydride
AgNO_3	Silver Nitrate
CDCl_3	Deuterated Chloroform
CH_3COOH	Acetic Acid
CoCl_2	Cobalt(II) Chloride

H.....	Hydrogen
H ⁺	Proton
H ₂ O	Water
H ₂ O ₂	Hydrogen Peroxide
HCl.....	Hydrochloric Acid
HO•	Hydroxy Radical
HOBr.....	Hypobromous Acid
HOCl.....	Hypochlorous Acid
HOO•	Hydroperoxy Radical
K ₃ Fe(CN) ₆	Potassium Ferricyanide
KCN	Potassium Cyanide
KHSO ₅	Potassium Peroxymonosulfate
MeOH-d ₄	Deuterated Methanol
NaOH	Sodium Hydroxide
Na ₂ IrCl ₆	Iridic Sodium Chloride
NEt ₃	Triethylamine
O ₂	Oxygen
O ₂ •-.....	Superoxide Radical Anion
PCl ₃	Phosphorous Trichloride

Chapter 1

Introduction and Background

1.1 Endogenous Exposome

The concept of the exposome was first introduced by Christopher Wild as a way to complement human genomics relative to investigating chronic diseases. The exposome can be defined as the culmination of environmental constituents the body comes in contact with, as well as the biological responses they elicit, from conception to death ¹. The definition was reintroduced in 2014 to include ‘endogenous processes’ to highlight the way the body reacts to environmental pressures and the complex chemistries resulting from the biochemical reactions that happen in our body like glycolysis and oxidative respiration ². Within the scope of the endogenous exposome, this project relates specifically to ribonucleic acids, or RNA.

RNA is transcribed from deoxyribonucleic acids (DNA) and in turn translates to proteins. Unlike DNA, RNA is usually single stranded and resides predominantly in the cytosol of the cell. Due to this, RNA is more susceptible to damage, primarily oxidative. RNA is also susceptible to alkylation, halogenation, and nitration. When RNA gets

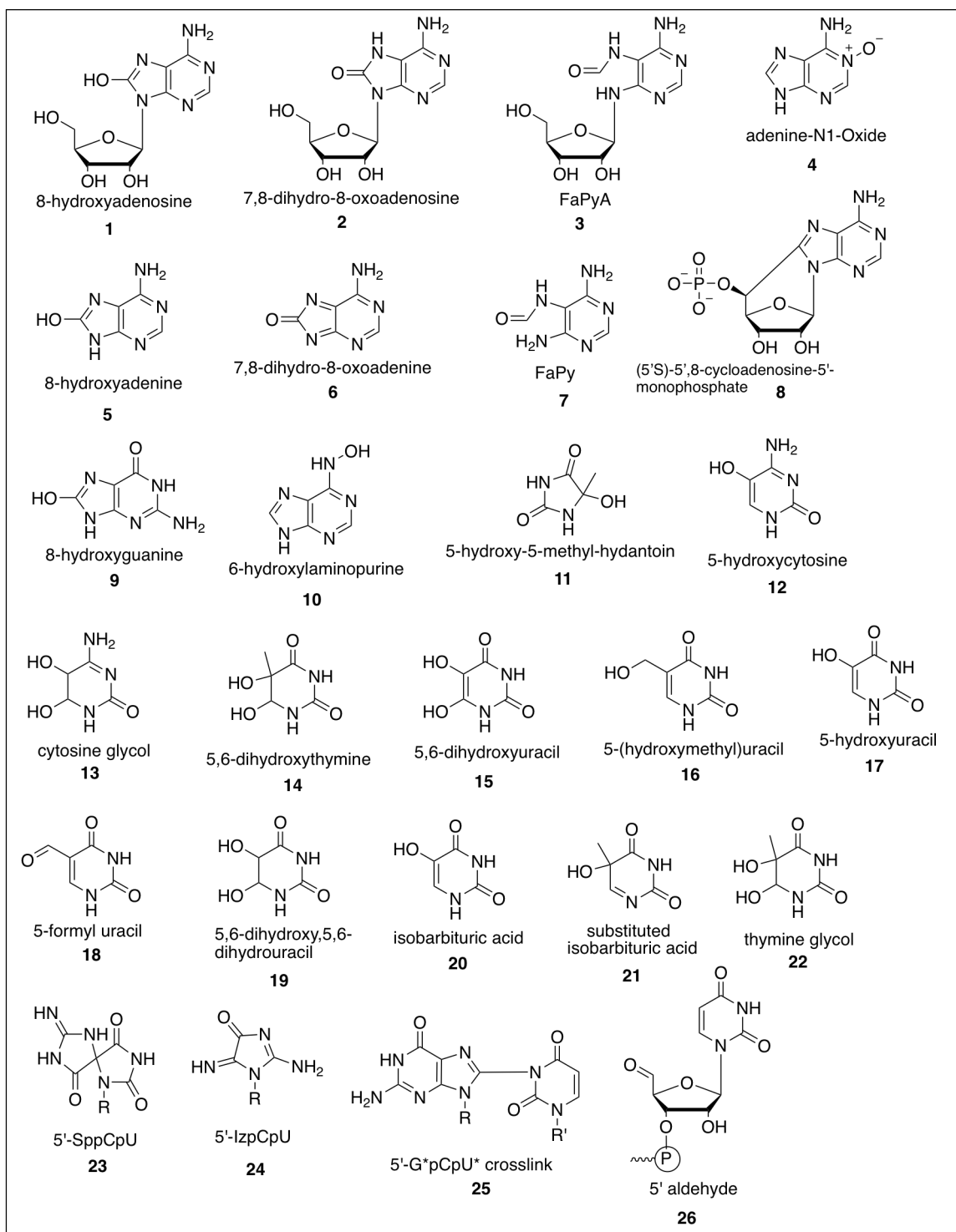


Figure 1-1. Hydroxyl Radical Lesions

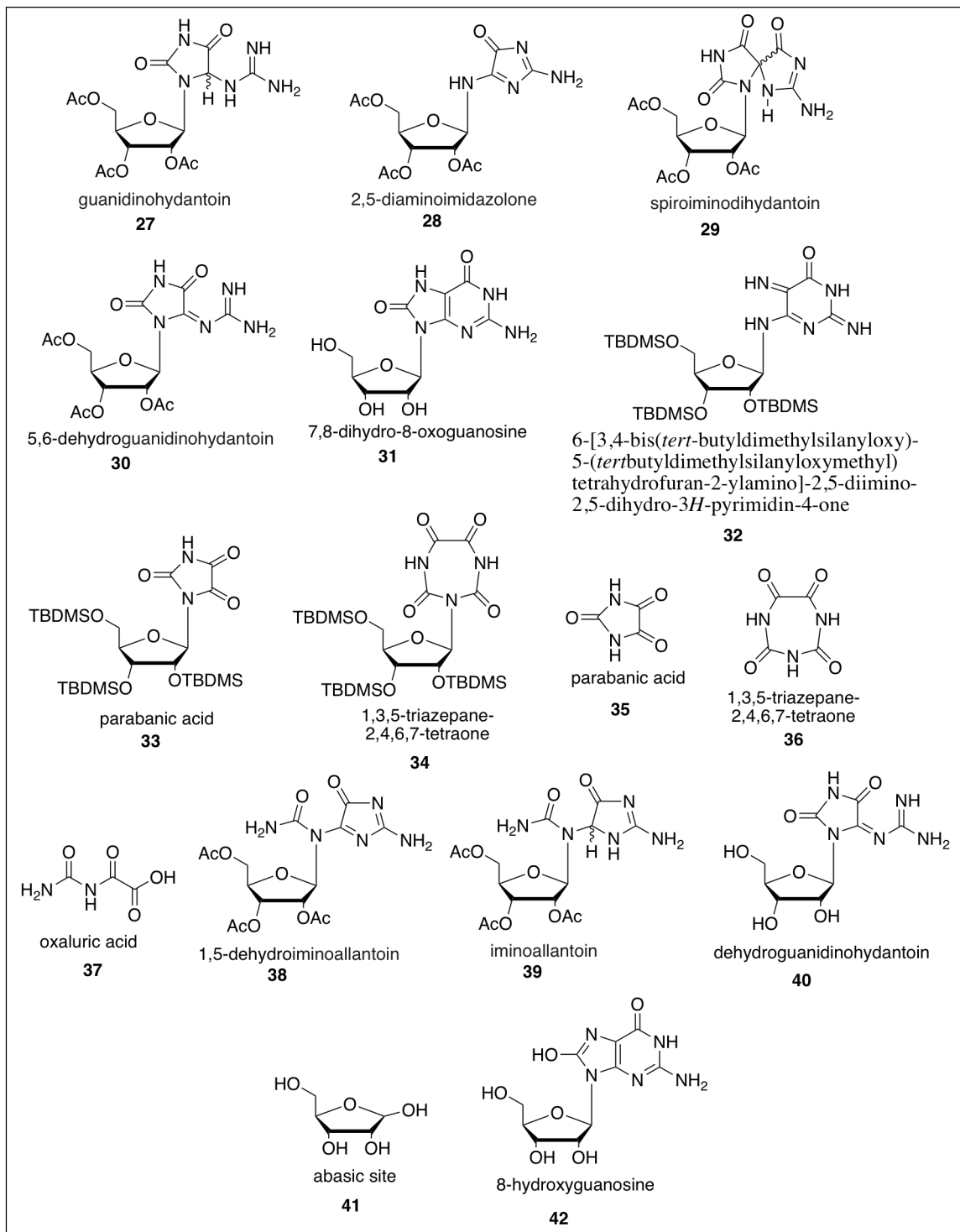


Figure 1-2. Singlet Oxygen Lesions

damaged, it can have deleterious ramifications. RNA lesions can lead to strand scission³⁻⁵, slow down or stop translation⁶⁻⁸, and change base pairing, making them mutagenic⁹⁻¹⁵. RNA lesions have been implicated in disease states, especially neurodegenerative diseases¹⁶⁻²⁹.

One of the most common damaging modifications that occurs in RNA is oxidation. The most common and promiscuous oxidative agent is the hydroxyl radical. The hydroxyl radical can abstract a hydrogen atom, undergo radical addition to the molecule, or perform a one-electron abstraction. Some lesions associated with this can be seen in **Figure 1-1**, **1-8** and **26**³⁰⁻³². However, the hydroxyl radical isn't the only oxidizing agent found in the body. Peroxyl radicals can also damage RNA as seen with compounds **5** and **9-16**³³. Lesions from the sulfoxyl radical are seen with compounds **17** and **18**³⁴. Compounds **19-22** are formed *via* the *tert*-butoxy radical³⁵. The phosphate radical anion also gives rise to **19** and **20** with substrates uracil and cytosine.³⁶ The carbonate radical anions yield **23-25** with a trinucleotide substrate³⁷.

Figure 1-2 shows lesions whose formation are associated with singlet oxygen, **27-36**³⁸⁻⁴⁰. Riboflavin as a one electron oxidant has been employed to study the product distribution of 8-oxoguanosine, wherein **27-30**, **35**, and **37-39** were the products⁴¹. Another method of investigating electron transfer-mediated damage involves the use of cationic manganese porphyrin associated with KHSO₅ as an oxidant⁴². The generated lesions were **29** and **40** when guanosine was in the anticodon of 17-mer tRNA (transfer RNA). This method of studying RNA oxidation also seems to be selective for G residues. Na₂IrCl₆, K₃Fe(CN)₆, and CoCl₂/ KHSO₅ are one electron oxidants used to study RNA

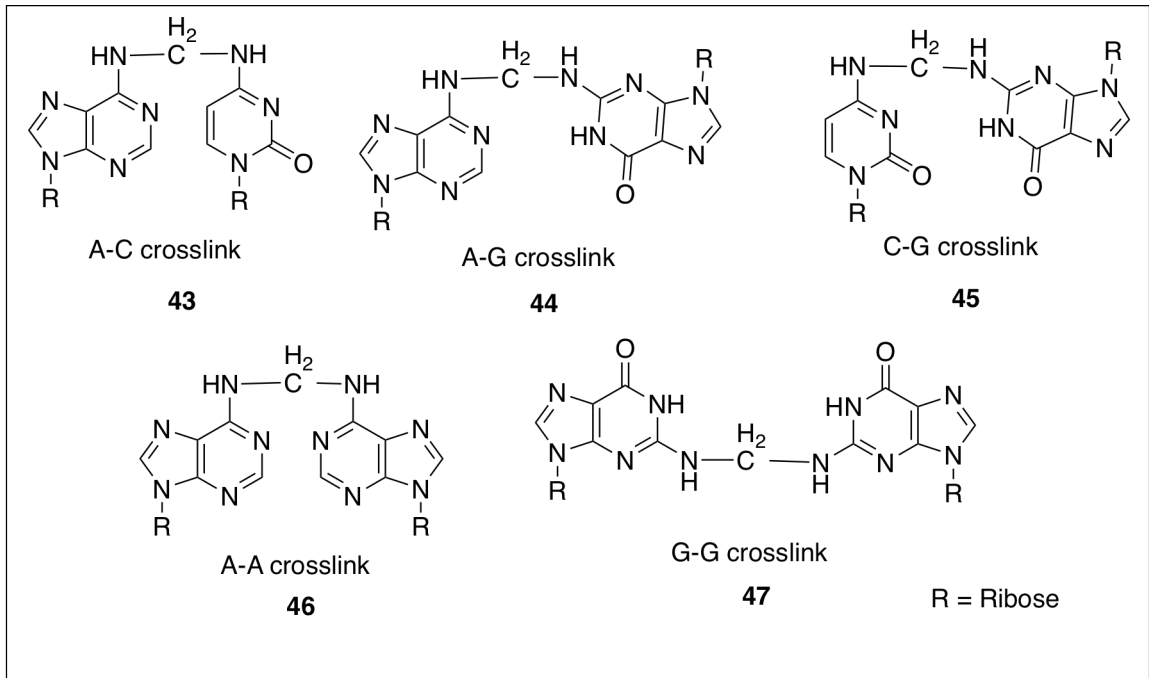


Figure 1-3. Formaldehyde Cross-Links

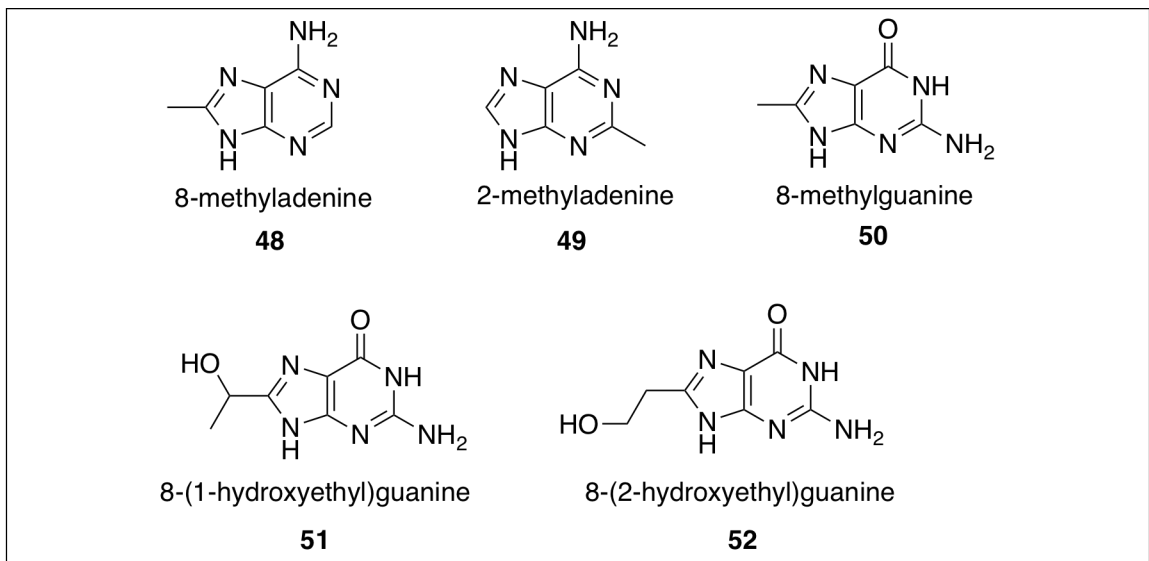


Figure 1-4. Alkylation Lesions

oxidation^{43,44}. Lesions formed in these systems are **27**, **32**, **38**, and **39**. Likewise, cytochrome c/H₂O₂ oxidizes guanosine in tRNA and leads to depurination to give **41** and **42**⁴⁵. Cytochrome c, which has also been observed to form a crosslink with RNA, could go on to trigger cellular apoptosis. Formaldehyde can also form cross-links with RNA, as seen in **Figure 1-3** with **43-47**⁴⁶.

Oxidative damage to RNA can have a multitude of biological effects. 8-Oxo-7,8-dihydroguanosine is the most common RNA oxidation lesion and is mutagenic^{9,12,47}. 8-Oxo-7,8-dihydroadenosine, 5-hydroxyuridine, and 5-hydroxycytosine have also been shown to be mutagenic, while 8-oxo-7,8-dihydroadenosine can destabilize RNA in a position-dependent manner^{11,13,14,48}. 5-Formyluridine may interfere with RNA metabolism and can base pair with guanine and cytosine, making it mutagenic as well¹⁰. Abasic sites in RNA templates follow the ‘A Rule’ as in DNA, where an adenosine is preferentially placed opposite the abasic site, however they are shown to be more stable and more promiscuous compared to their DNA counterpart^{49,50}. Some lesions have been studied for their effect on mRNA (messenger RNA) translation. Abasic sites are non-coding lesions that interrupt peptide synthesis. 8-Oxo-7,8-dihydroguanosine, 8-oxo-7,8-dihydroadenosine, 5-hydroxyuridine, and 5-hydroxycytidine lead to a mixture of full-length and truncated peptides⁸.

Another form of RNA damage is alkylation. Normally, the body specifically methylates using SAM (S-Adenosyl Methionine). However, agents, such as the methyl radical, can methylate RNA as well. The methyl radicals usually add to the base, as seen in **Figure 1-4**, **48-50** thus disrupting base pairing⁵¹. Methylation of tRNAs has been

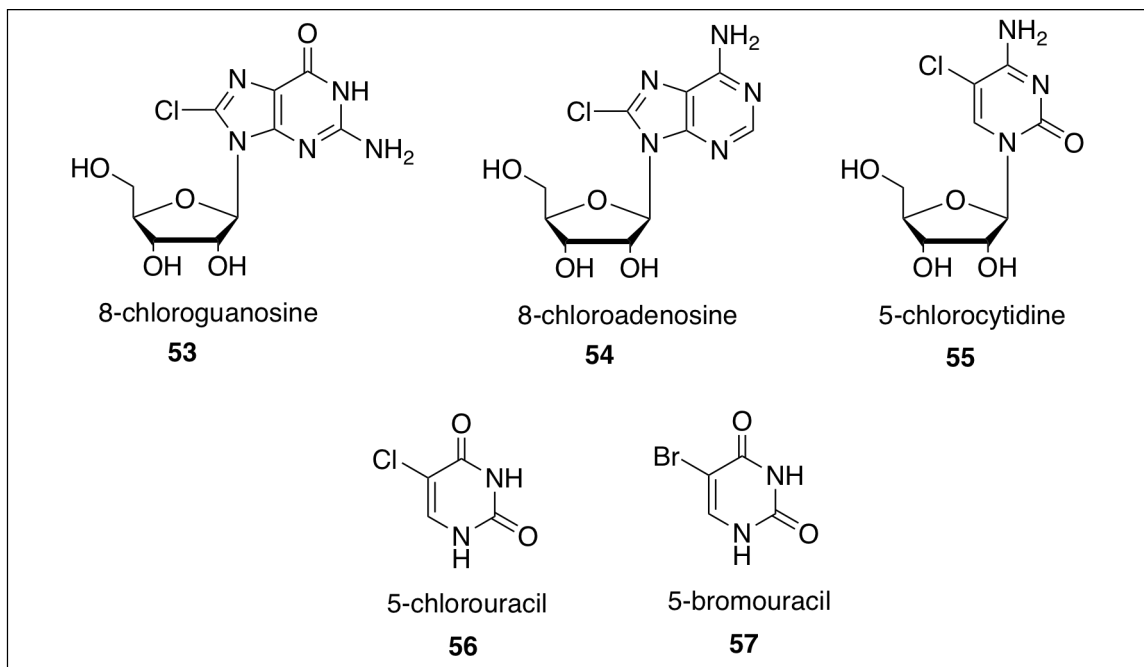


Figure 1-5. Halogenation Lesions

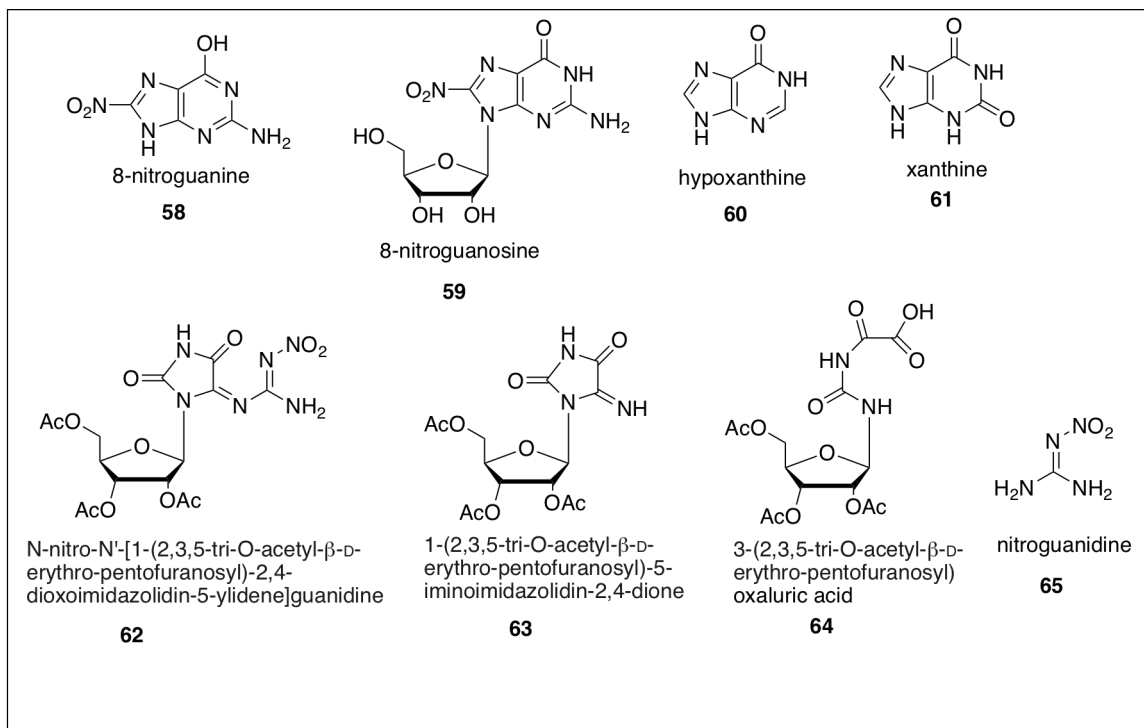


Figure 1-6. Nitration Lesions

shown to inhibit aminoacylation, while methylation of both tRNAs as well as mRNAs can inhibit translation⁵². Methylation can also interfere with tRNA – ribosome binding at the A site⁵³. In a Fenton system, ethanol oxidation can lead to oxidation/alkylation lesions **51** and **52**^{54, 55}. 8-Methylguanosine, as well as 8-oxo-7,8-dihydroguanosine, have been shown to induce cell proliferation in certain cells, but more work has to be done to investigate the potential of these lesions in free radical-mediated carcinogenesis⁵⁶.

RNA can also be halogenated as seen in **Figure 1-5**, and has been shown to be more susceptible to chlorination than DNA⁵⁷. Lesions associated with this can be seen with hypochlorous acid (HOCl), **53-56**⁵⁸⁻⁶⁰. In an eosinophil peroxidase system, uracil was brominated to **57** by HOBr⁶¹. In pre-mRNA, 5-chloro and 5-bromouridine were unable to be recognized for splicing⁶². 8-Chloroadenosine is currently in phase I clinical trials for the treatment of chronic lymphocytic leukemia and has also been looked at for the treatment of breast cancer^{63, 64}. The mechanisms by which it acts are inhibition of global transcription and reduction of intracellular ATP (adenosine triphosphate) concentrations.

Figure 1-6 shows lesions associated with nitration, another important type of damage that affects RNA. Peroxynitrite can react with guanine to form 8-nitroguanine, **58**⁶⁵. In an *H. pylori* infection study, it was shown that there was an increase in 8-nitroguanine due to inflammation⁶⁶. This was also correlated to an increase in gastric cancer in patients. 8-Nitroguanosine, **59**, has been shown to be involved in the pathogenesis of murine pneumotropic virus infections⁶⁷. Unlike DNA, 8-

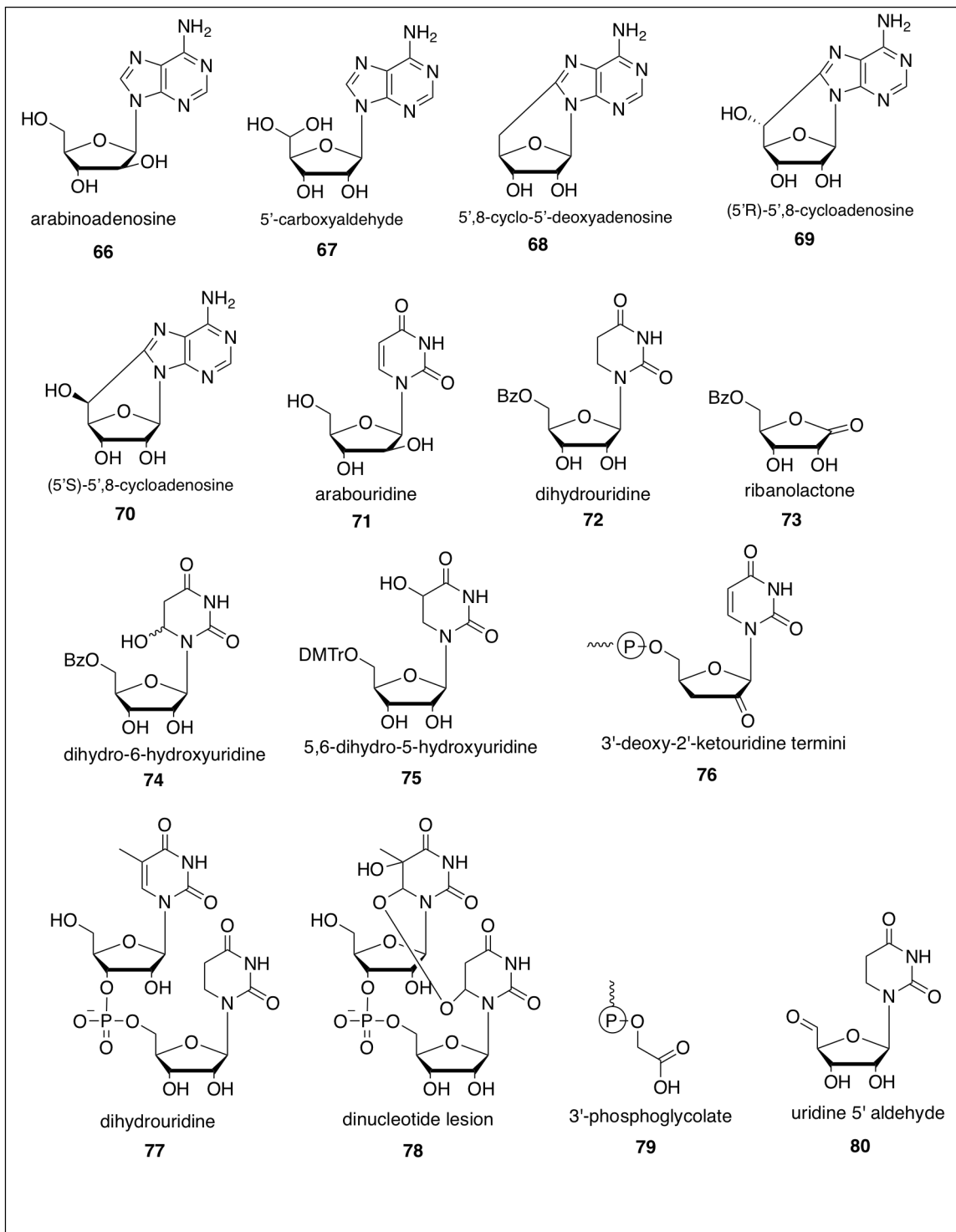


Figure 1-7. Chemical Elucidation Lesions

nitroguanosine is very stable in RNA⁶⁸. Nitric oxide can cause the formation of **60** and **61** from deamination of guanine and adenine in calf thymus RNA. In rat kidneys, hypoxanthine was shown to induce lipid peroxidation^{69, 70}. When ribose-protected 8-oxo-7,8-dihydroguanosine was exposed to peroxyxynitrite, **62-65** were formed⁷¹.

The endogenous exposome has also been investigated using organic chemistry. Not only has the automated synthesis of oligonucleotides been invaluable, but so has the use of radical precursors. There have been a few groups to utilize this method, including our own, to identify **66-80** in **Figure 1-7**^{3-5, 72-77}. Dihydrouridine is the most common post-transcriptional modification in tRNA and has the opposite effect of a pseudouridine or ribose methylation modification⁷⁸. Dihydrouridine stabilizes the C2'-endo sugar conformation, giving the RNA more flexibility. This may have an undesired destabilizing effect when present.

1.2 Organic Chemistry as a Tool in the Study of RNA Damage

1.2.1 Radical Precursors

Radical precursors have been a useful tool for the selective generation of a radical at a site of interest to mimic oxidative damage. This is accomplished by employing a photolabile group, such as a pivaloyl moiety. This functionality when subjected to UV wavelengths ≥ 240 nm causes a Norrish Type I cleavage as seen in **Figure 1-8**⁷⁹.

Giese et al. were the first to employ these radical precursors to study RNA when they looked at the 4' position of the sugar in adenosine⁷⁷. A radical precursor with a

photolabile acetyl group was incorporated into a strand of RNA then irradiated to generate the 4' radical. Comparison with the corresponding radical in DNA revealed that the cleavage of the latter was 3 times faster than that of the RNA. Under aerobic conditions it was also found that the presence of the 2' hydroxyl influenced product composition by slowing a β -elimination of **84** allowing for the attack of water leading to the formation of the 3'-phosphoglycolate **79** as seen in **Scheme 1-1**. The 2' radical in the adenosine monomer was also generated and found to result in the release of adenine⁷².

Greenberg et al. group has also worked with radical precursors in RNA. First, their group focused on the C5 and C6 positions of the base in uridine, looking at both monomer and oligomer reactive intermediates. When generating the 5,6-dihydrouridin-6-yl radical in the monomer, dihydro-6-hydroxyuridine **74** was the major product found under aerobic conditions and under anaerobic conditions the reduced dihydrouridine **72** was the major product³. It was also reported that the reactivity was similar to the corresponding radical in DNA. The same radical in an oligomer led to strand scission through the formation of a C2' radical intermediate, as did the C5 dihydrouridine radical, which was faster^{4, 5, 75}. The proposed mechanism is shown in **Scheme 1-2** leading to the formation of the 3'-deoxy-2'-ketouridine termini **76**. They also generated the C2' radical in uridine and found that it underwent base release to give uracil in monomer studies⁷⁴. Oxygen and β -mercaptoethanol were unable to compete with base loss while thiol was able to compete in less polar solvents. Greenberg's group then incorporated their C2' radical precursor into strands of RNA and investigated its reactivity^{73, 80}. They found that C2'-hydrogen atom abstraction was the rate-determining step, and, like before,

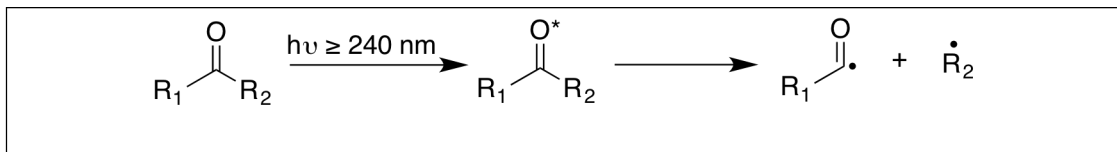
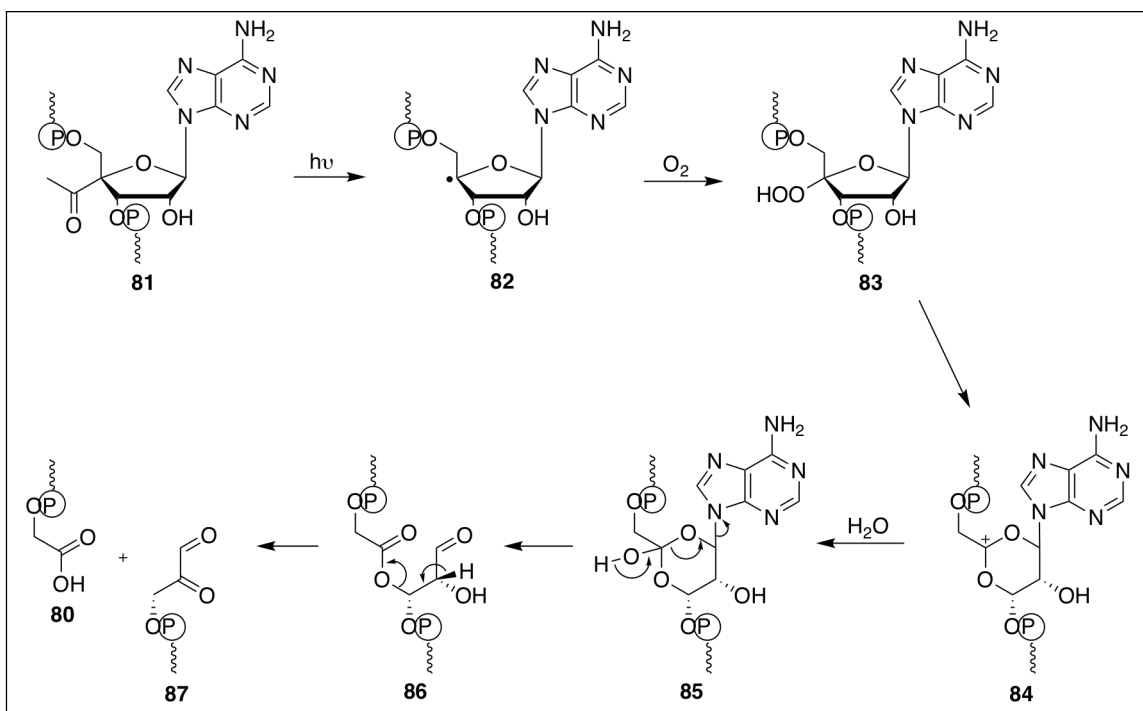


Figure 1-8. Norrish Type I Cleavage

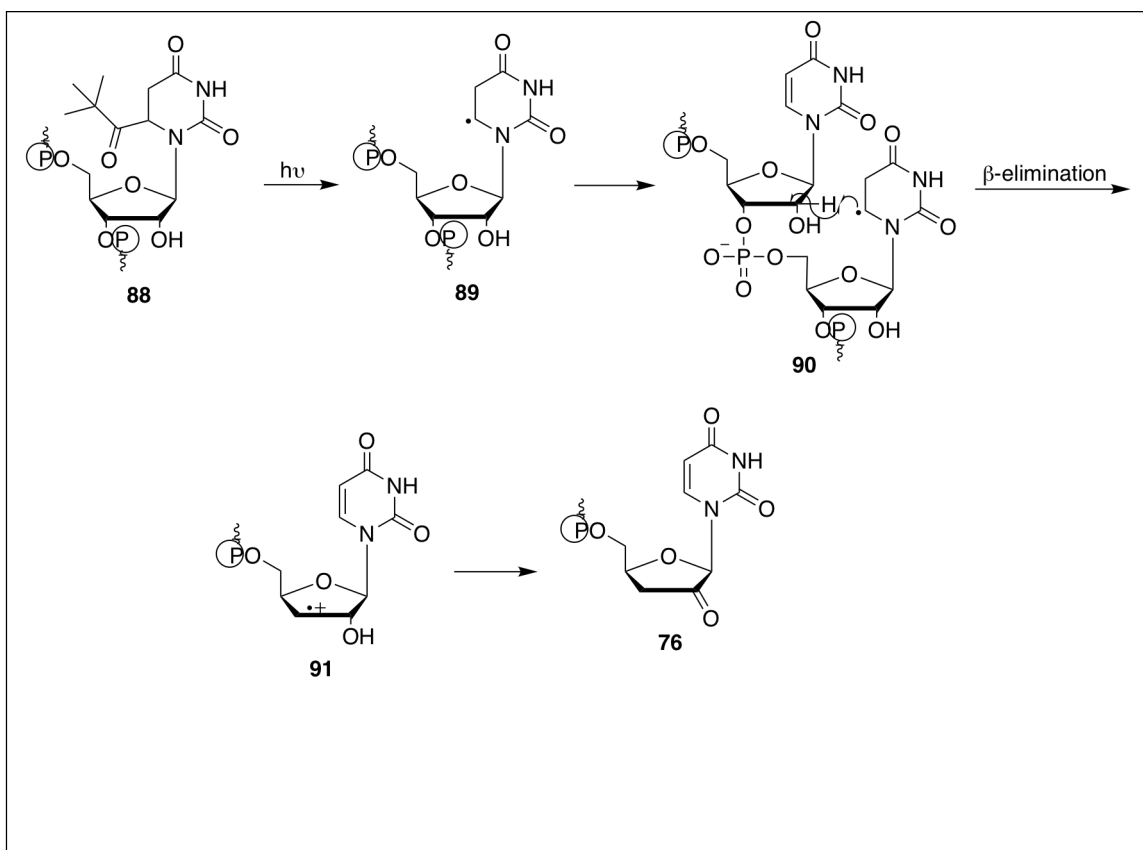


Scheme 1-1. Adenosyl 4' Radical Mechanism

the major product was the 3'-deoxy-2'-ketouridine termini **76**, as well as a 3'-phosphate derived from the site where strand scission occurred. This happened with or without oxygen in the presence of β -mercaptoethanol. Uracil loss still occurred in modified strands, however its formation was unable to compete with the elimination of the 3'-phosphate leading to strand scission. Due to the major group of reactive intermediates formed on pyrimidines, the hydroxyl radical adducts, and the role the C2'-hydroxyl plays in the rate limiting step, it can be suggested that RNA appears to be more susceptible to strand scission than DNA.

1.2.2 Oligoribonucleotide Synthesis

Automated oligonucleotide synthesis is widely used to synthesize strands of DNA and RNA for various uses. The basis for which this works is the coupling of the phosphorous moiety of the added nucleotide with the free hydroxyl group of the extending strand of nucleotides. This process includes coupling, oxidation, and deprotection steps, which are repeated until the desired length is achieved. Automated RNA synthesis can be accomplished using a variety of building blocks, however the most commonly used are the phosphoramidites and H-phosphonates. Both utilize the same base protecting groups such as benzoyl and isobutyryl and both use similar sugar protecting groups such as dimethoxytrityl for 5' protection and a *tert*-butyldimethylsilyl for 2' protection. The difference is in the phosphorus moiety at the 3' position, which both utilize P(III). Many phosphoramidites contain a bulky diisopropyl-amino group and require a cyanoethyl protecting group while the H-phosphonate contains a phosphoryl



Scheme 1-2. 5,6-Dihydrouridin-6-yl Radical Mechanism

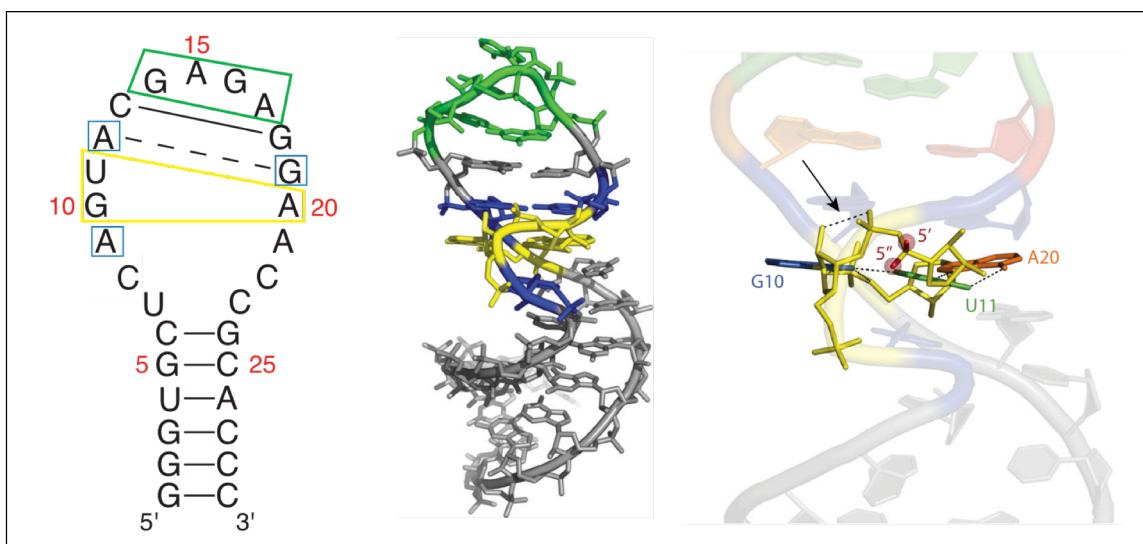


Figure 1-9. Sarcin/Ricin Loop RNA ³²

functionality as well as a hydrogen atom attached to the phosphorous. Due to this difference, the automated synthesis using these methods must be altered slightly. H-Phosphonates have a tetrahedral geometry like P(V) compounds and lack a pair of electrons, making them electrophilic⁸¹. This makes them much more stable to oxidation compared to phosphoramidites, which require an oxidation step after each coupling cycle. Aside from this, the steps for the two are very similar. First, the dimethoxytrityl group is removed giving a free 5' hydroxyl to couple with the added monomer, which is facilitated by an activator. After the condensation, there is a capping step to ensure no further reactivity. Finally, in the case of the phosphoramidite, there is the oxidation step. This is repeated until the oligomer is of the desired length, then the final base and 2' deprotections, as well as the removal of the solid support, are performed.

In vitro transcription has also been useful for the investigation of RNA damage. This method utilizes an RNA polymerase and a template strand of DNA, which transcribes to the complementary strand of RNA. This method was utilized by Tullius et al. to obtain the sarcin/ricin loop RNA as shown in **Figure 1-9**. It was determined that the structural environment that occurs within a GUA triplet in RNA leads to unusual reactivity of a hydroxyl radical with the C5' hydrogen of the uridine³². They showed that the hydroxyl radical abstracted the C5' hydrogen on U11 in the sarcin/ricin loop and gave rise to the 5' aldehyde terminus **26** at the cleavage site as shown in **Figure 1-10**.

All of the work reported in this chapter was done with RNA substrates. However, much of the work in the field of nucleic acid damage has been conducted on DNA. Though the presence of the 2' hydroxyl group changes its reactivity, many of the lesions found in DNA may also be present in RNA. More work must be completed in this field in

order to further elucidate the diversity of RNA damage and to be able to better compare the damage that occurs in nucleic acids.

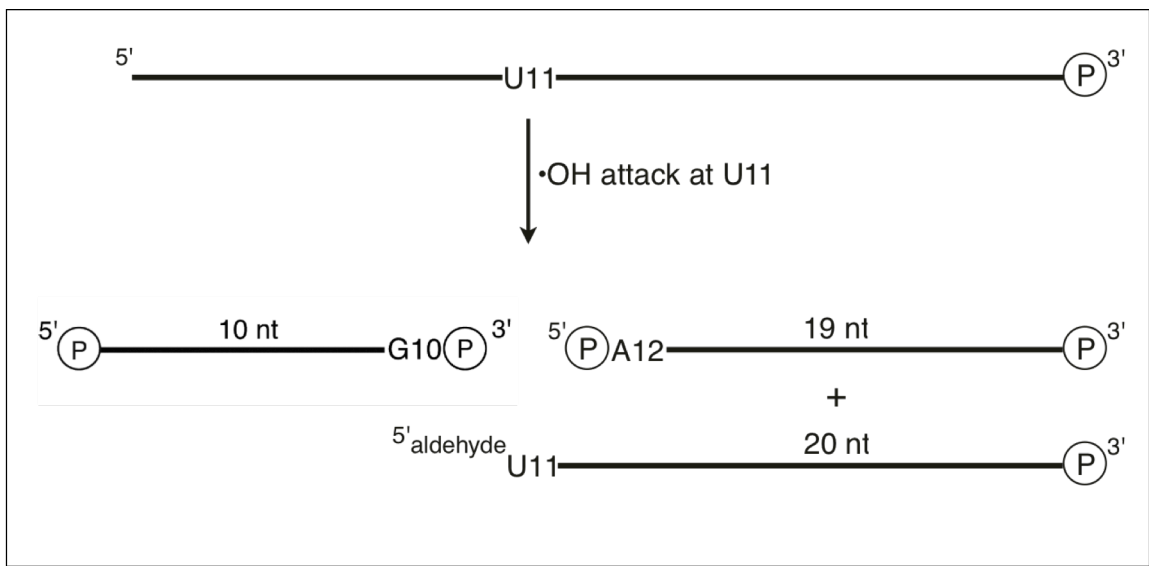


Figure 1-10. Formation of 5' aldehyde³²

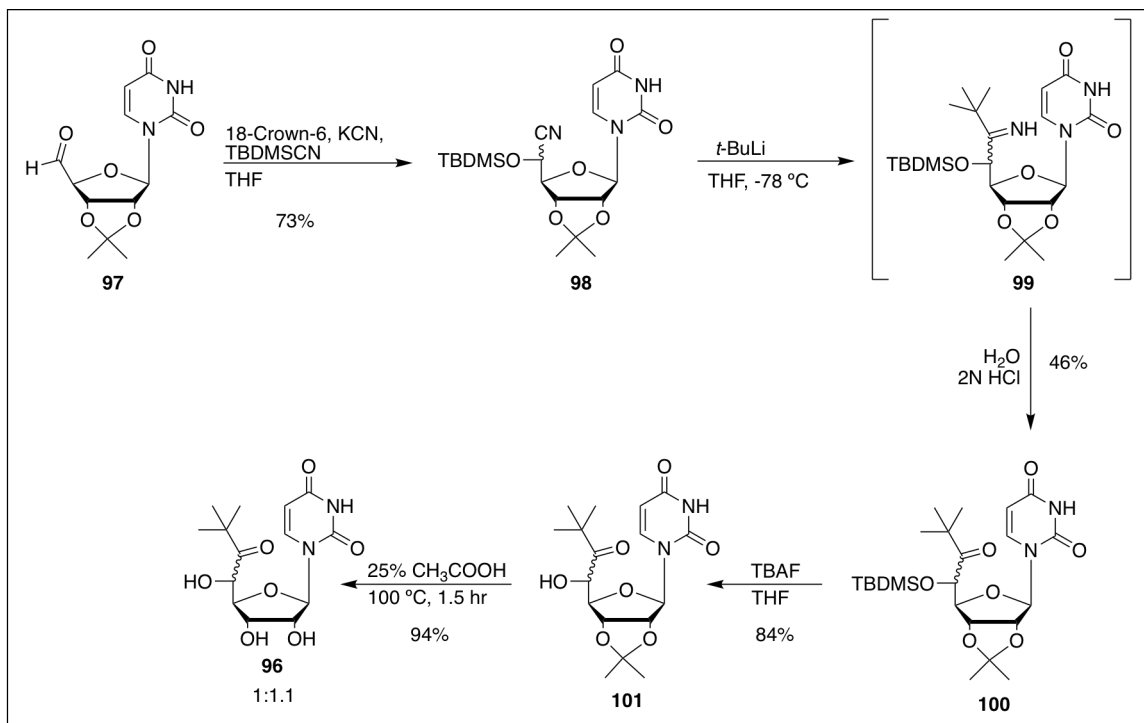
Chapter 2

Results and Discussion

2.1 5' Uridinyl Radical Precursor Synthesis

As shown in Chapter 1, RNA damage can take many forms and have a multitude of adverse effects in the body. Both the base and sugar portion of RNA are susceptible, and there are many different causative agents. Of these, oxidation is the most prevalent, and the most promiscuous oxidizing agent is the hydroxyl radical. To study oxidative damage, it is helpful to look at the reactivity of specific nucleic acid radicals. Just as Giese's group investigated the 4' position of adenosine and Greenberg's group the 2' position of uridine, so has our group studied the 5' position of uridine, which Tullius' group has shown to generate the radical of interest under certain RNA structural motifs in the presence of the hydroxyl radical ^{32, 72-74, 76, 77, 80}.

To site-specifically generate the 5' radical, a *tert*-butyl ketone, or pivaloyl group, was installed on the 5' position of the sugar in uridine. This was previously accomplished by our group and the first part of this project was to validate and optimize the synthesis shown in **Scheme 2-1** ⁷⁶. Conditions explored for the optimization can be shown in **Table 2-1**.



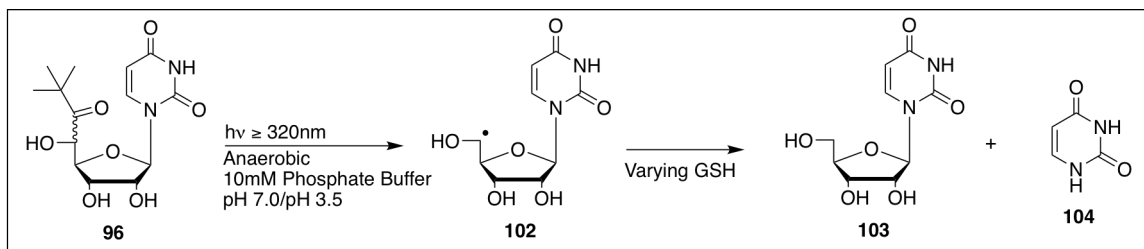
Scheme 2-1. Synthesis of 5' Uridinyl Radical Precursor

<i>t</i> -BuLi Reaction Conditions	Crude % Yield of 99	Hydrolysis Conditions	% Yield of 100
15 min	-	THF, pH 7	15
2 min	43	-	-
4 min	71	-	-
8 min	63	-	-
4 min	-	THF, pH 6	30
4 min	-	pH 6, added ACN, 30 °C	18
4 min	-	ACN, pH 4, 60 °C	46

Table 2-1. 5' Uridinyl Radical Precursor Synthetic Optimization

Compound **97** was made from uridine following standard literature protocols, with the entire synthesis itself being based on work by Chatgililoglu⁸²⁻⁸⁴. Compound **97** was treated with *tert*-butyldimethylsilyl cyanide (TBDMSCN) in the presence of catalytic amounts of 18-Crown-6 and potassium cyanide (KCN) in THF (tetrahydrofuran) to give compound **98** as a mixture of diastereomers⁷⁶. Cyanohydrin **98** was then converted to the imine using *t*-BuLi (*tert*-butyllithium) at – 78 °C followed by hydrolysis to give protected ketone **100**⁷⁶. This was followed by the use of TBAF (tetrabutylammonium fluoride) to remove the 5' silyl group and then acid hydrolysis to give radical precursor **96**⁷⁶.

Table 2-1 shows the conditions used in the optimization of the reaction conditions leading from cyanohydrin **98** to ketone **100**. The first step was to find the best reaction time for the *t*-BuLi reaction. Initially, 15 min was used, however it should not take so long for the completion of this reaction and prolonging the exposure to *t*-BuLi may lower the yield. *t*-BuLi is an extremely reactive reagent, so much so that it will combust upon exposure to the air. It will even react with the solvent, THF⁸⁵. Using 4 equivalents of *t*-BuLi means that there is extra reagent present able to participate in side reactions after it has reacted with the nucleoside at the nitrile. Starting from 2 min, the time was increased to 4 min then 8 min, which showed that 4 min had the highest crude yield for the reaction. Next, the hydrolysis conditions were tested. Initially THF was used with water and 2 N HCl (hydrochloric acid), however when the pH was tested, it was still too close to 7. In order to facilitate the acid hydrolysis without the removal of any of the acid labile protecting groups, the pH was maintained between 5-6. Simply reducing the *t*-BuLi reaction time to 4 min and the pH to 6 doubled the percent yield, but the reaction still



Scheme 2-2. Preliminary Anaerobic 5' Uridinyl Radical Formation

pH	GSH (mM)	Uridine 103 (%)	Uracil 104 (%)
7.0	0	n.d.	33.9 ± 0.83
	2	7.12 ± 0.12	43.7 ± 4.98
	4	9.44 ± 0.02	44.3 ± 3.26
	8	7.30 ± 1.45	47.8 ± 2.86
	16	7.79 ± 2.96	43.1 ± 3.75
	24	10.5 ± 0.87	42.3 ± 5.51
	32	15.4 ± 0.55	43.3 ± 3.21
3.5	0	n.d.	13.6 ± 1.15
	2	30.3 ± 0.75	10.4 ± 0.45
	4	45.7 ± 2.10	11.2 ± 1.09
	8	54.6 ± 0.45	11.2 ± 0.17
	16	66.9 ± 2.39	10.7 ± 0.17
	24	70.0 ± 0.72	8.38 ± 1.52
	32	75.9 ± 1.82	8.55 ± 1.26

Table 2-2. Preliminary Anaerobic 5' Uridinyl Radical Product Distribution

took 91 hours. On the second attempt, just water and acid were used while adjusting the temperature to 30 °C after 23 hours. There were solubility issues with the nucleoside, and, after 43 hours, ACN was added. The initial, and continued heating, as well as the addition of acetonitrile both appeared to drive the reaction forward. However, the total reaction time was 94 hours. Acetonitrile (ACN) is a more polar aprotic solvent than THF and can aid in the solubility of the compound, which was seen. Going forward, ACN and heat were added from the beginning, heating up to 60 °C. The pH was also adjusted more acidic, to pH 4. This reaction only took 90 min to complete with much higher yield.

2.2 5' Uridinyl Radical Precursor Photolysis

To investigate the reactivity of the 5' uridinyl radical, precursor **96** was photolyzed under a variety of conditions. Our lab previously ran a study showing the differences in product distribution when photolyses were performed at either pH 7.0 or pH 3.5 under anaerobic conditions with varying amounts of GSH for one hour, which is shown in **Scheme 2-2** and **Table 2-2**⁷⁶. This showed that uracil formation was not GSH dependent like uridine formation. It also showed that reduction by GSH was much more efficient at lower pH, which confirms what has been reported in the literature⁸⁶.

The 5' uridinyl radical reactivity was further investigated in my project under anaerobic conditions using a physiologically relevant amount of GSH (8 mM) in phosphate buffer at pH 7.0 with varying photolysis times (1, 2, 4, 8, 15, 30, and 60 min). This was done to ascertain whether any reactive intermediates were forming within that 60 min window.

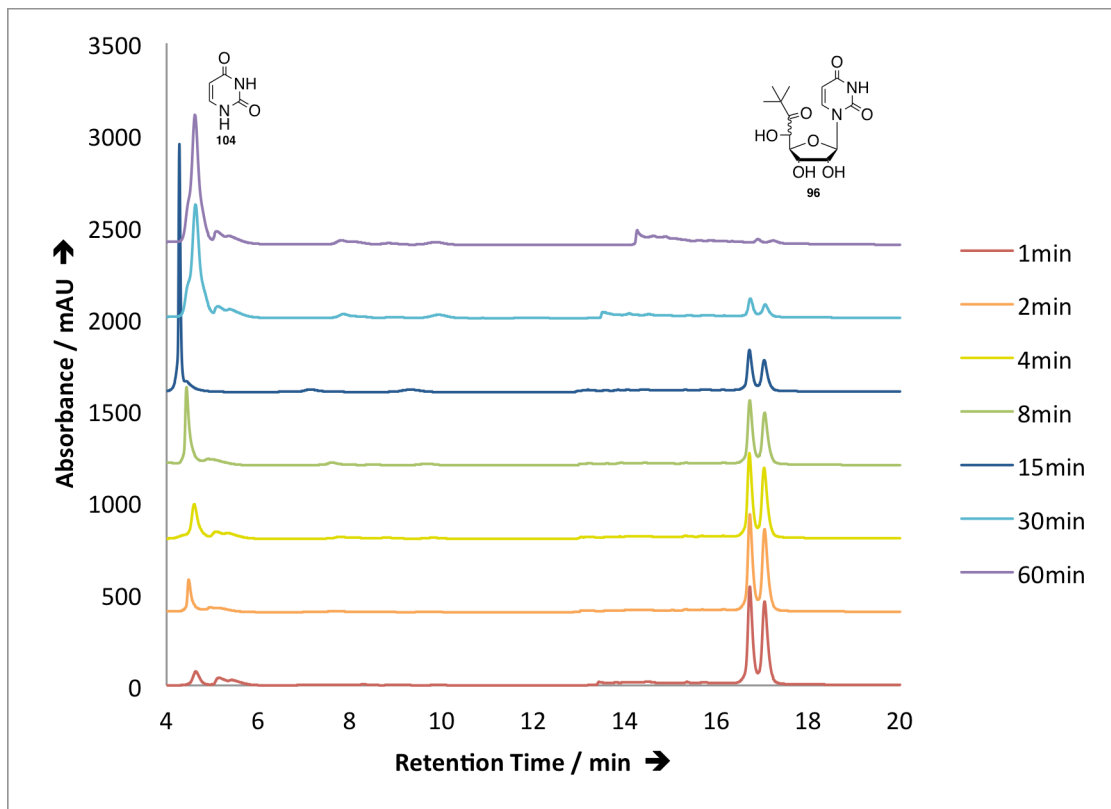


Figure 2-1. RP-HPLC Results for Anaerobic Photolysis of 5' Uridinylyl Radical Precursor

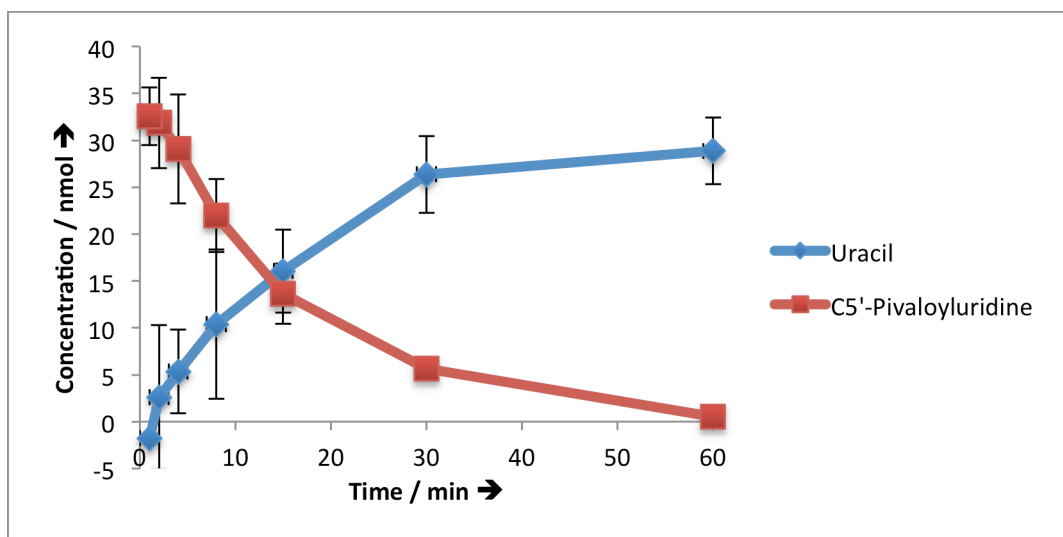


Figure 2-2. Conversion of 5' Uridinylyl Radical Precursor to Uracil under Anaerobic Conditions

The RP-HPLC results were overlaid in **Figure 2-1**. Here, there were no UV active intermediates found and uridine was below the limit of detection. The mass recovery in this study was 80 – 93% and it was shown that after 60 min there was a 78% conversion of radical precursor **96** to uracil, as seen in **Figure 2-2**.

This project was continued to photochemically generate the 5' uridiny radical under aerobic conditions using 8 mM GSH and 10 mM ammonium acetate at pH 7.0 and 3.5 for 30 min. The results can be seen in **Figure 2-3**. It was found that the 5' aldehyde (11.8 min), or its corresponding hydrate, as well as the base elimination product uracil (12.9 min) were the major products formed. Small amounts of uridine (18.8 min) were also observed, as well as some peaks that we were unable to identify at both pH 7 and 3.5. The peak at 24 min was seen when GSH was injected alone, and the masses observed are 613.5 and 307.4 in positive ion mode and 611.4 in negative ion mode. This points towards oxidized GSH (GSSG) at 612.63. The peaks at 39.2 min and 43.9 min are in the blank for the column. The percent conversion of the precursor to radical related products was 56.2% at pH 7 and 58.0% at pH 3.5. Of the amount of radical generated, there was 1.7% conversion to uridine at pH 7 and pH 3.5. For uracil, there was 14.4% conversion at pH 7 and 15.1% at pH 3.5. Based on calculations for the 5' aldehyde, it is 108.7% and 110.0% for pH 7 and 3.5 respectively. For the 5' aldehyde, the calculations are slightly skewed due to what must be another UV active compound with a very similar retention time. Accounting for uracil and uridine formation, as well as a small amount of unknown product, the conversion to the 5' aldehyde is close to 80%. This is also confirmed with what is seen on MS.

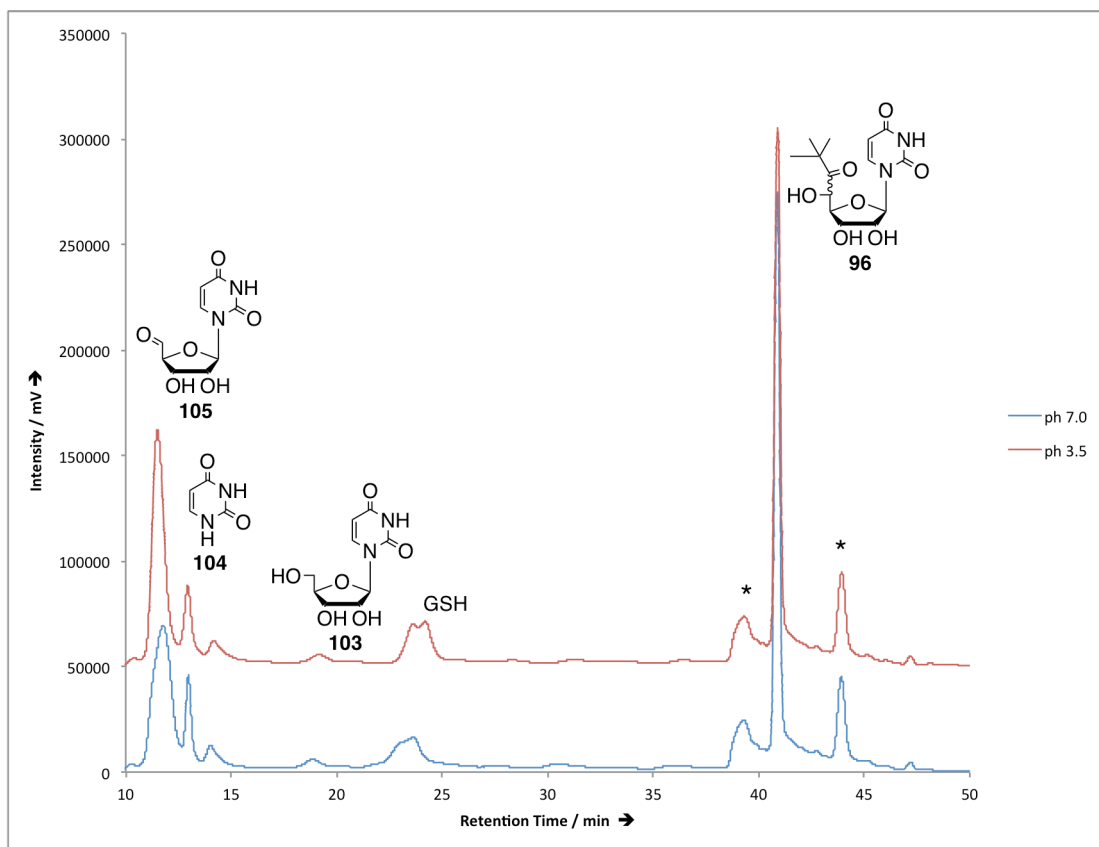
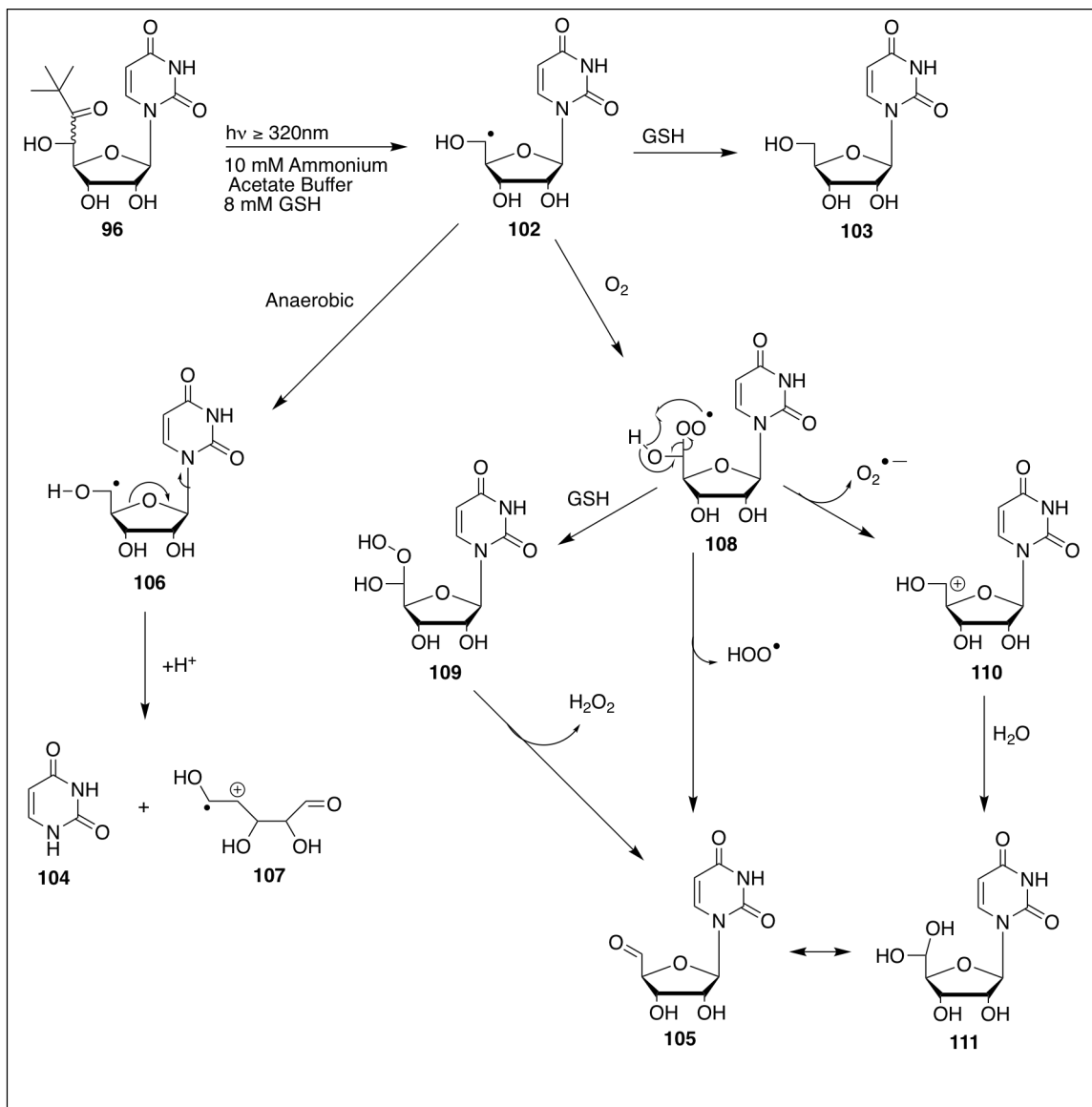


Figure 2-3. RP-HPLC Results for Aerobic Photolysis of 5' Uridiny Radical Precursor
 * Column Contaminants



Scheme 2-3. Proposed Mechanism of 5' Uridinyl Radical

2.3 5' Uridinyl Radical Mechanism

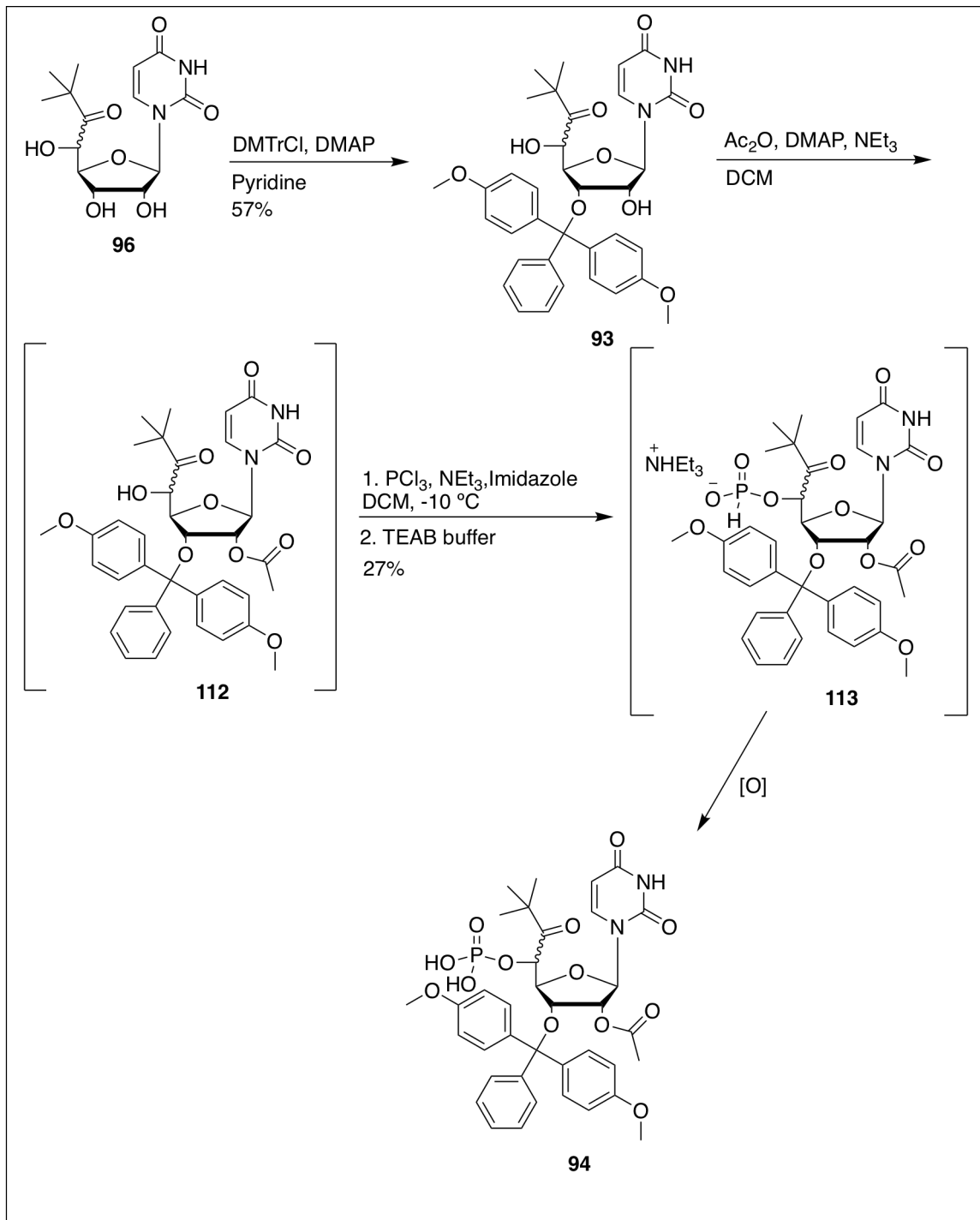
The formation of uridine suggests that radical **102** is indeed generated and involved in the degradation of the molecule. Uracil formation was found to be independent of glutathione concentration under anaerobic conditions, and there were no other compounds identified. This suggests a competition, under anaerobic conditions, of base release and reduction by GSH, the former of which appears to be much faster under physiological conditions. **Scheme 2-3** shows the proposed mechanism under both anaerobic and aerobic conditions ⁷⁶. It is suggested that the C4'-O bond breaks in a β -elimination reaction to release uracil, facilitating ring opening. β -Elimination reactions are common when dealing with radicals, and uracil is an excellent leaving group ^{87, 88}. This mechanism appears to be in competition with the fate of the radical under aerobic conditions. Oxygen can react at a rate to of 10^9 , while GSH reacts at a rate of 10^6 in nucleic acid radicals, which explains why so little uridine is seen ⁸⁹.

Once intermediate **108** is formed, there are three possible outcomes. First, reduction by GSH can lead to the hydroperoxide **109**, which would have a similar reaction rate as before. Second is the loss of $O_2^{\bullet-}$, superoxide radical anion, leaving a positive charge on the 5' position, which would lead to the nucleophilic addition of water and hydrate formation. This process can occur at a rate of 10^4 making it unable to compete with GSH ⁹⁰. The third possibility is the loss of HOO^{\bullet} , hydroperoxy radical. This loss has been shown to occur via the mechanism shown in **Scheme 2-3** ⁹¹⁻⁹⁶. It is also a much more favorable outcome compared to the loss of $O_2^{\bullet-}$ or the H atom abstraction event that could occur ^{91, 93}. A study of α -amino-peroxyl radicals showed that

in a vacuum, the H atom abstraction event was on the order of 10^8 while the loss of the HOO• was 10^{12} . They also found that α -hydroxy-peroxyl radicals were more likely to undergo loss of the HOO• and that a polar solvent system made it even more favorable⁹¹. MS and RP-HPLC confirmed formation of **105** as the major lesion, and, to our knowledge, this is the first time this lesion has been identified in a monomeric ribonucleoside.

2.4 5' Uridinyl Radical Precursor H-Phosphonate Synthesis

In order to further study the reactivity of the 5' uridinyl radical and the effect of RNA damage, the 5' uridinyl radical precursor **96** needs to be incorporated into strands of RNA. To accomplish this, **96** was derivatized to contain a 5' H-phosphonate with the appropriate protecting groups for RNA synthesis. The synthesis is shown in **Scheme 2-4**. First, **96** was protected with a dimethoxytrityl group on the 3' hydroxyl using DMTrCl and DMAP in pyridine. Compound **93** was then acylated at the 2' hydroxyl using Ac₂O (acetic anhydride) NEt₃ and a catalytic amount of DMAP in DCM^{97,98}. This was then subjected to H-phosphonate addition conditions using PCl₃ and NEt₃, then TEAB. Compound **113** was confirmed by HRMS, however the ³¹P NMR clearly shows that it was oxidized to monophosphate **94**. Though H-phosphonates are more stable to oxidation than their phosphoramidite counterparts, it can still occur⁹⁹. Likely, the purification of compound **113** left it too exposed. In the future, this should be used without further purification, as are many H-phosphonates, to prevent this issue.



Scheme 2-4. Synthesis of Uridine 5' H-Phosphonate Radical Precursor

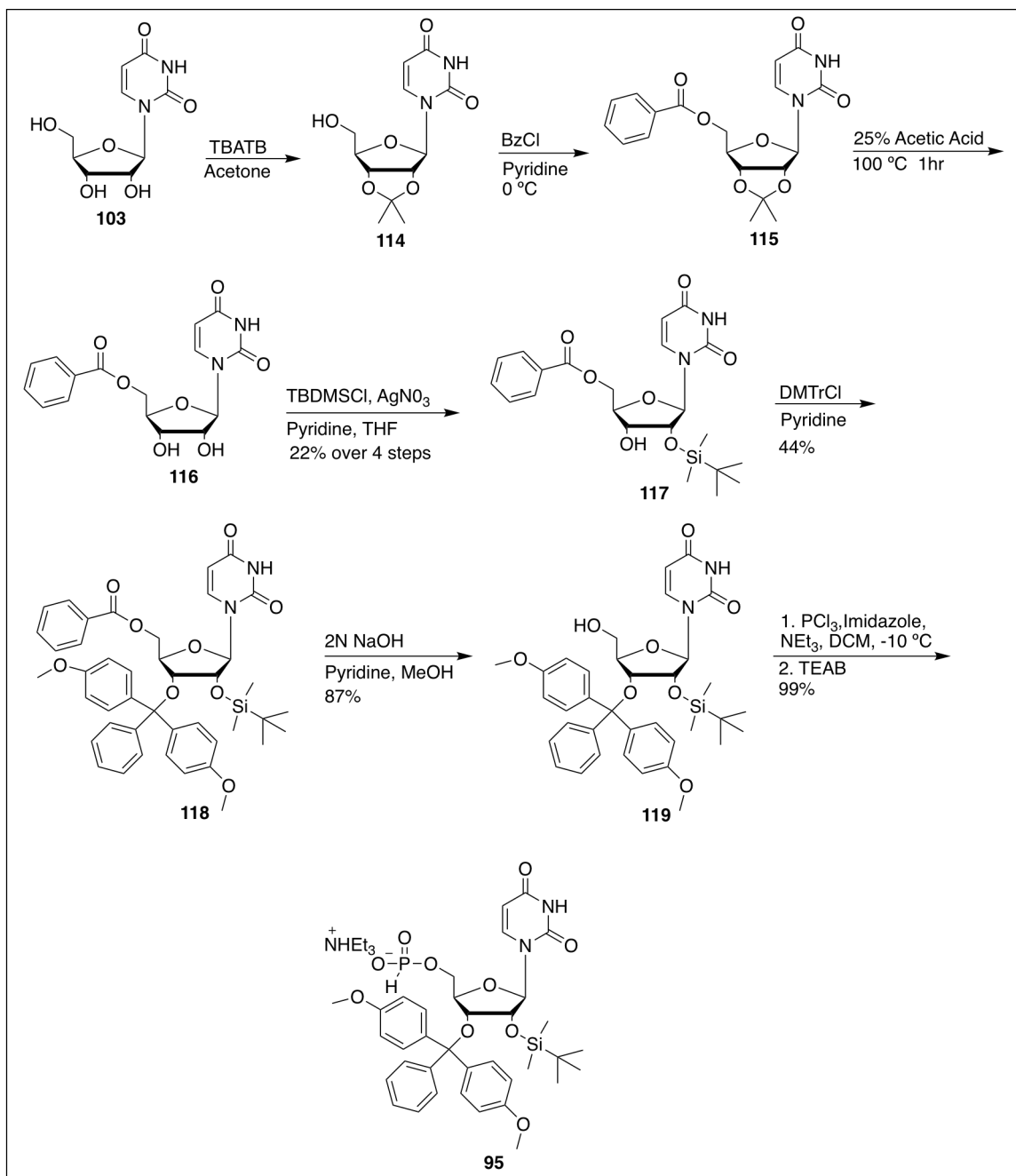
2.5 H-Phosphonate Monomer Syntheses

2.5.1 Overview of 5' H-Phosphonate Synthesis

In order to incorporate our radical precursor into strands of RNA, we endeavored to design a new automated RNA synthesis technique based on the assembly of the oligomer in the 5' to 3' direction using 5' H-phosphonates. This included the synthesis of the four RNA building blocks, uridine, cytidine, adenosine, and guanosine 5' H-phosphonates. Recently, there has been work to develop RNA synthetic methods in the reverse direction, from 5' to 3', using 5' phosphoramidites¹⁰⁰⁻¹⁰³. These methods still utilize a 2' O-silyl protecting group with a dimethoxytrityl protecting group at the 3' position. This work was foundational for our studies.

Synthesis in the 5' to 3' direction would allow for the introduction of a variety of 3' modifications such as ligands and chromophores to aid research in multiple fields. For example, this could be applicable for RNA therapeutics by providing easy access to 3' modifications on the sense strand of siRNA, which should not interfere with recognition since antisense siRNA strands guide target recognition¹⁰⁰. Synthesis in the 5' to 3' direction also allows for more steric modifications on the sugar moiety, especially the 5' position when using an H-phosphonate, suiting our purposes well.

There are also advantages to using H-phosphonate chemistries over that of phosphoramidites. Phosphoramidites require an oxidation step after each coupling cycle,



Scheme 2-5. Synthesis of Uridine 5' H-Phosphonate

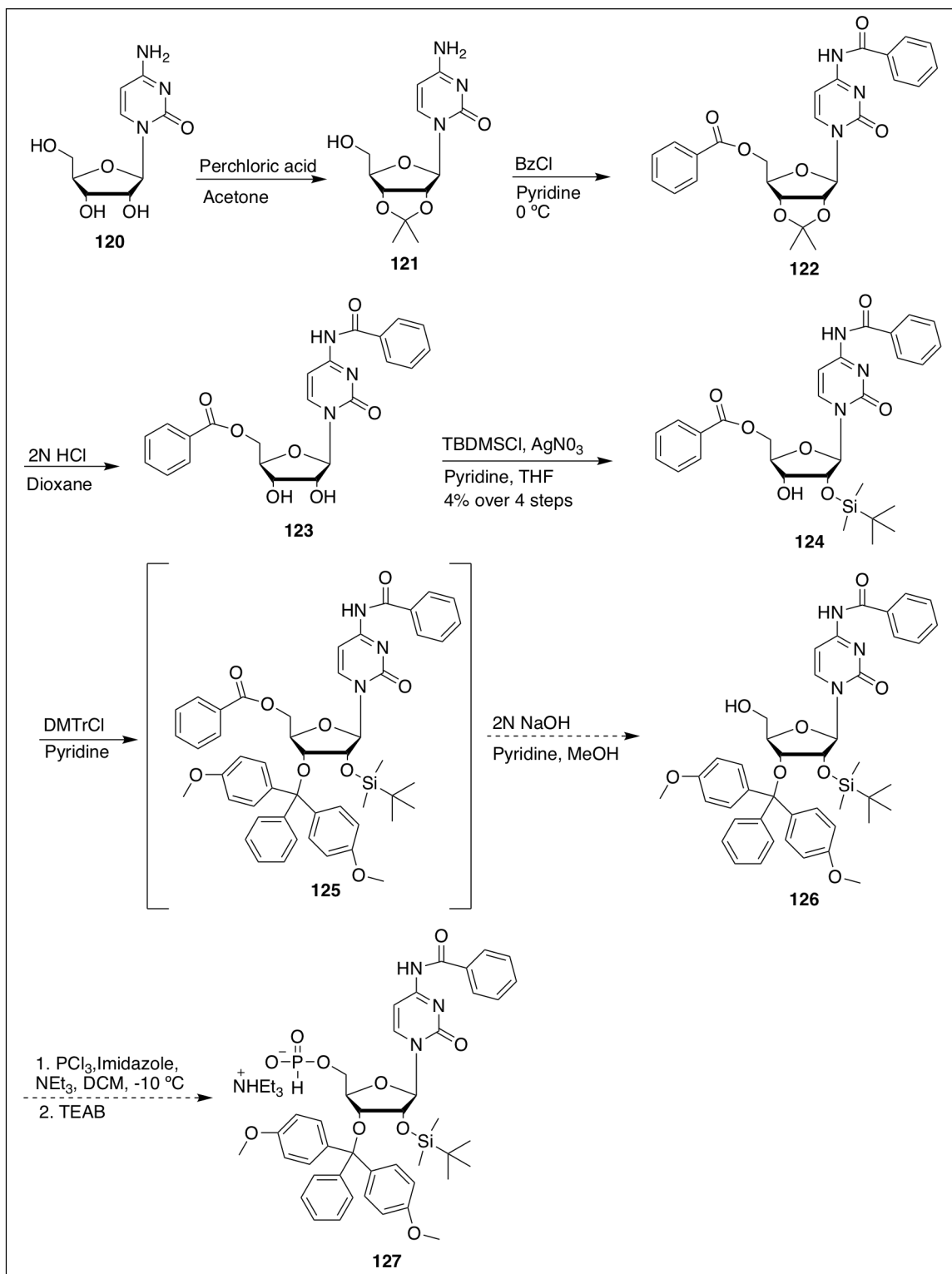
whereas in H-phosphonate chemistry only one oxidation step is required at the end of oligomer assembly. This cuts down the number of steps required in each cycle, as well as the cycle time. H-phosphonates are also more stable than phosphoramidites and have the potential to be recovered after coupling^{99,104}. H-phosphonate chemistry also uses some of the same readily available reagents used in the phosphoramidite approach.

2.5.2 Uridine 5' H-Phosphonate Synthesis

The synthesis of **95** can be seen in **Scheme 2-5**. First, uridine was protected using a catalytic amount of tetrabutylammonium tribromide (TBATB) in acetone to give **114**⁸³. This was benzoylated at the 5' position using benzoyl chloride (BzCl) in pyridine to afford **115**¹⁰⁵. Fully protected **115** then underwent acidic hydrolysis using 25% acetic acid and heat to give **116**⁷⁶. Compound **116** was selectively silylated at the 2' position using *tert*-butyldimethylsilyl chloride and silver nitrate (AgNO₃) in a mixture of pyridine and tetrahydrofuran¹⁰⁶. Compound **117** was then protected using dimethoxytrityl chloride in pyridine to give **118**, which was finally selectively deprotected using 2N sodium hydroxide (NaOH) in pyridine and methanol to give 5' hydroxyl **119**¹⁰². The addition of the H-phosphonate was completed using phosphorous trichloride, imidazole, and triethylamine in dichloromethane using protocols developed in our lab¹⁰⁷.

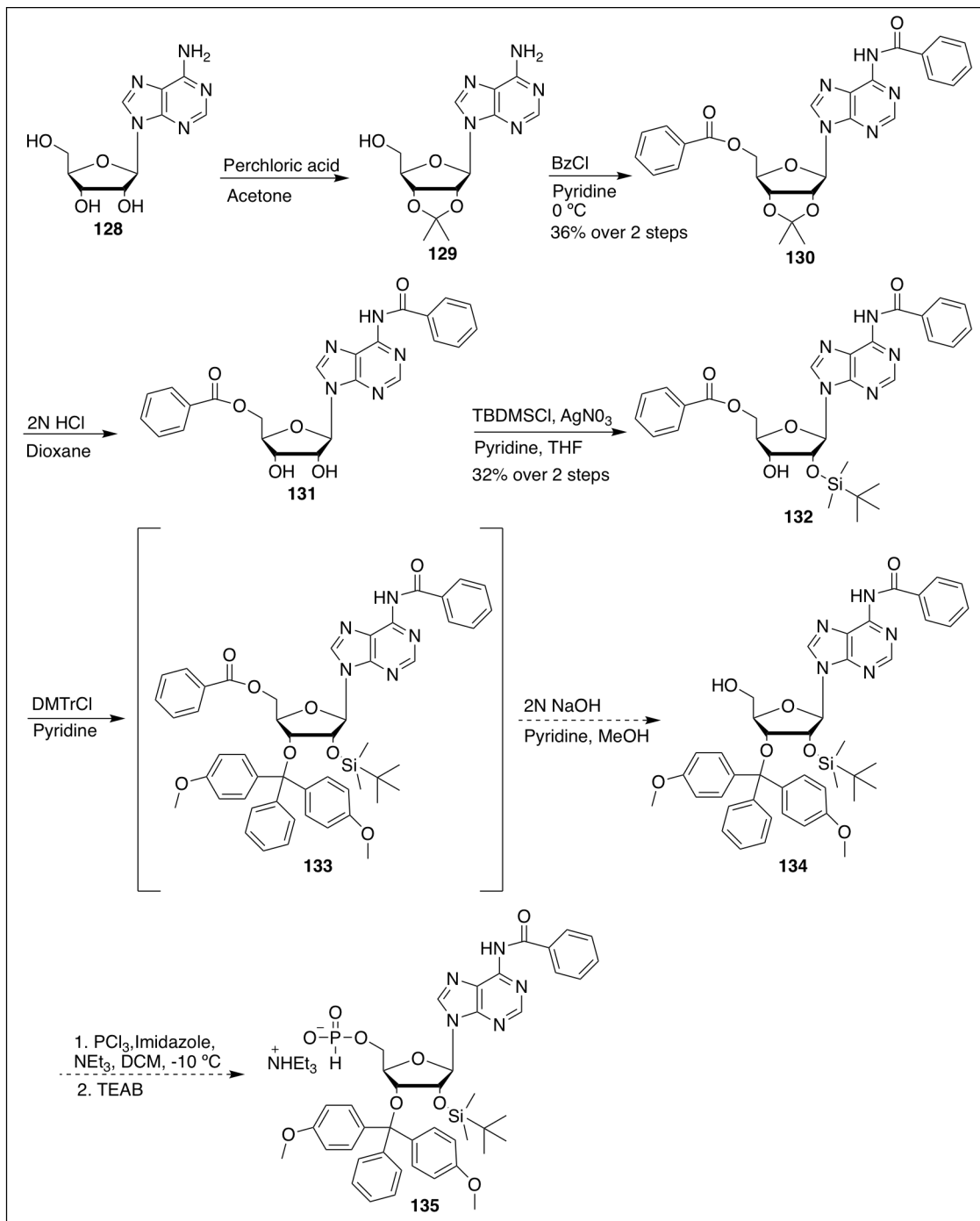
2.5.2 Cytidine 5' H-Phosphonate Synthesis

The attempted synthesis of **127** can be seen in **Scheme 2-6**. First, cytidine was protected using a catalytic amount of perchloric acid in acetone to give **121**¹⁰⁸. This was benzoylated at the 5' position, as well as on the exocyclic amine, using benzoyl chloride



Scheme 2-6. Attempted Synthesis of Cytidine 5' H-Phosphonate

(BzCl) in pyridine to afford **122**¹⁰⁵. These first two steps had high crude yields and likely are not part of the problem of the low combined yield. Fully protected **122** then underwent acidic hydrolysis using dilute hydrochloric acid (HCl) in dioxane to give **123**¹⁰². This gave a crude yield of approximately 50%. In the workup, the crude product is redissolved in EA and THF, but there are solubility issues. Even when DCM and MeOH are added to help this, it is possible that in the aqueous washes some product is lost. This acid hydrolysis has also been shown to be problematic with the guanosine and adenosine nucleosides, perhaps causing more compound degradation than initially thought. To further investigate this, **122** should be purified before the acid hydrolysis. Compound **123** was selectively silylated at the 2' position using *tert*-butyldimethylsilyl chloride and silver nitrate (AgNO₃) in a mixture of pyridine and tetrahydrofuran¹⁰⁶. The selective protection gives preference to the formation of the desired 2' hydroxyl, however it also gives rise to the protected 3' hydroxyl and bis-protected compounds. Combined, these steps delivered **124** in a very low yield. In part, this was due to the lack of purification in between any of these reactions. In the future, compound **122** should be purified before moving forward, and, most importantly, compound **123** should be very carefully purified so as to give one pure compound going into the selective protection. Another possibility is to deploy a scheme similar to what was adopted with guanosine where there is a transient protection of the three hydroxyl groups to start with a base protected nucleoside, removing the need to perform the isopropylidene removal. Compound **124** was then protected using dimethoxytrityl chloride in pyridine to give **125** in a crude yield of 84%, after which an attempt to selectively deprotect by using 2N sodium hydroxide (NaOH) in pyridine and methanol failed¹⁰². In the future, **125** should be purified before moving forward to ensure



Scheme 2-7. Attempted Synthesis of Adenosine 5' H-Phosphonate

there is one pure product to test the reaction with.

2.6.2 Adenosine 5' H-Phosphonate Synthesis

The attempted synthesis of **135** can be seen in **Scheme 2-7**. First, adenosine was protected using a catalytic amount of perchloric acid in acetone to give **129**¹⁰⁸. This was benzoylated at the 5' position, as well as on the exocyclic amine, using benzoyl chloride (BzCl) in pyridine to afford **130**¹⁰⁵. Here, it is also possible to form a compound with two Bz protecting groups on the exocyclic amine. It was observed that this had started to form before the starting material was fully consumed, so the reaction was stopped. In the future, varying conditions should be tested to optimize this procedure, with the possibility of moving forward with an adenosine compound obtaining three Bz protecting groups. Fully protected **130** then underwent acidic hydrolysis using dilute hydrochloric acid in dioxane to give **131**¹⁰². Again, this saw low crude yields. Compound **131** was selectively silylated at the 2' position using *tert*-butyldimethylsilyl chloride and silver nitrate (AgNO₃) in a mixture of pyridine and tetrahydrofuran¹⁰⁶. Compound **132** was then protected using dimethoxytrityl chloride in pyridine to give **133** with a crude yield of 88%, which was pushed forward to attempt the selective deprotection using 2N sodium hydroxide (NaOH) in pyridine and methanol which failed¹⁰². In the future, **133** should be purified before going forward to ensure a single, pure product to test the reaction with. As stated before, the scheme could be modified as in the case of guanosine to remove the need to use an isopropylidene protecting group on the 2' and 3' hydroxyls, which should reduce the number of steps used and potentially increase the overall yields.

2.7.2 Guanosine 5' H-Phosphonate Synthesis

An appropriately protected guanosine is required for RNA synthesis and multiple candidates were investigated in an attempt to achieve this (**Figure 2-4**). The protecting groups could vary for the base, however the isobutyryl (Ibu) group is the most commonly used for guanosine, likely due to its easier removal during automated synthesis. For the 5' protecting group, the benzoyl (Bz) group aided greatly in compound solubility.

First, the same approach as used in the other nucleosides was attempted, seen in **Scheme 2-8**. Guanosine was protected using a catalytic amount of perchloric acid in acetone to give **137**¹⁰⁸. This was benzoylated at the 5' position, as well as on the exocyclic amine, using benzoyl chloride (BzCl) in pyridine to afford **138**¹⁰⁵. Fully protected **138** then underwent acidic hydrolysis using dilute hydrochloric acid in dioxane to give **139**¹⁰². This reaction was performed multiple times, and the product would sometimes degrade during the reaction. However, more success was observed during faster acidic hydrolysis prompted by heat. Compound **139** was selectively silylated at the 2' position using *tert*-butyldimethylsilylchloride and silver nitrate in a mixture of pyridine and tetrahydrofuran¹⁰⁶. This reaction was more successful when the previous reaction was purified or the product was washed with a saturated sodium bicarbonate solution and brine, however **139** does have some solubility issues. Under these reaction conditions, the compound was also prone to decompose via base release. Compound **140** was then protected using dimethoxytrityl chloride in pyridine to give **141**¹⁰². The selective

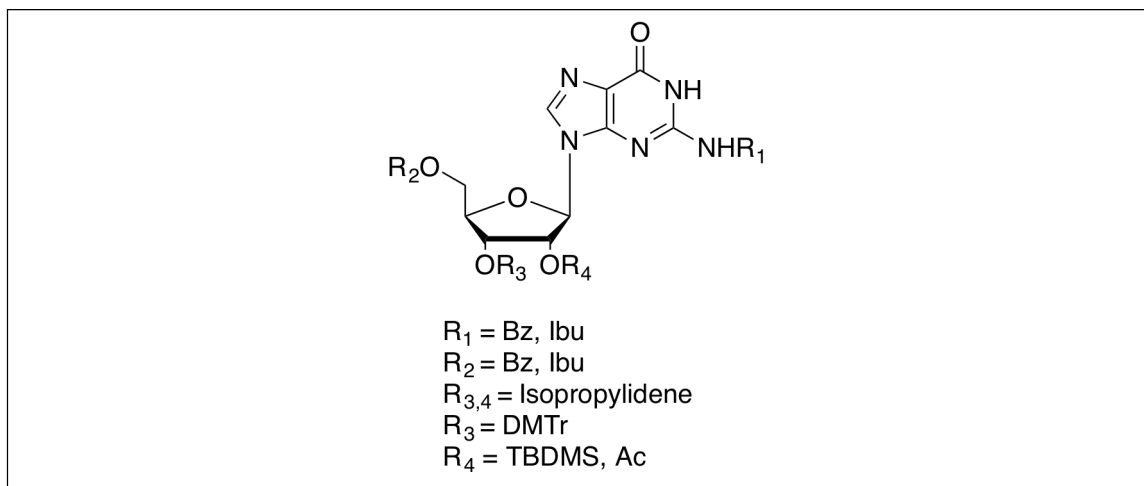
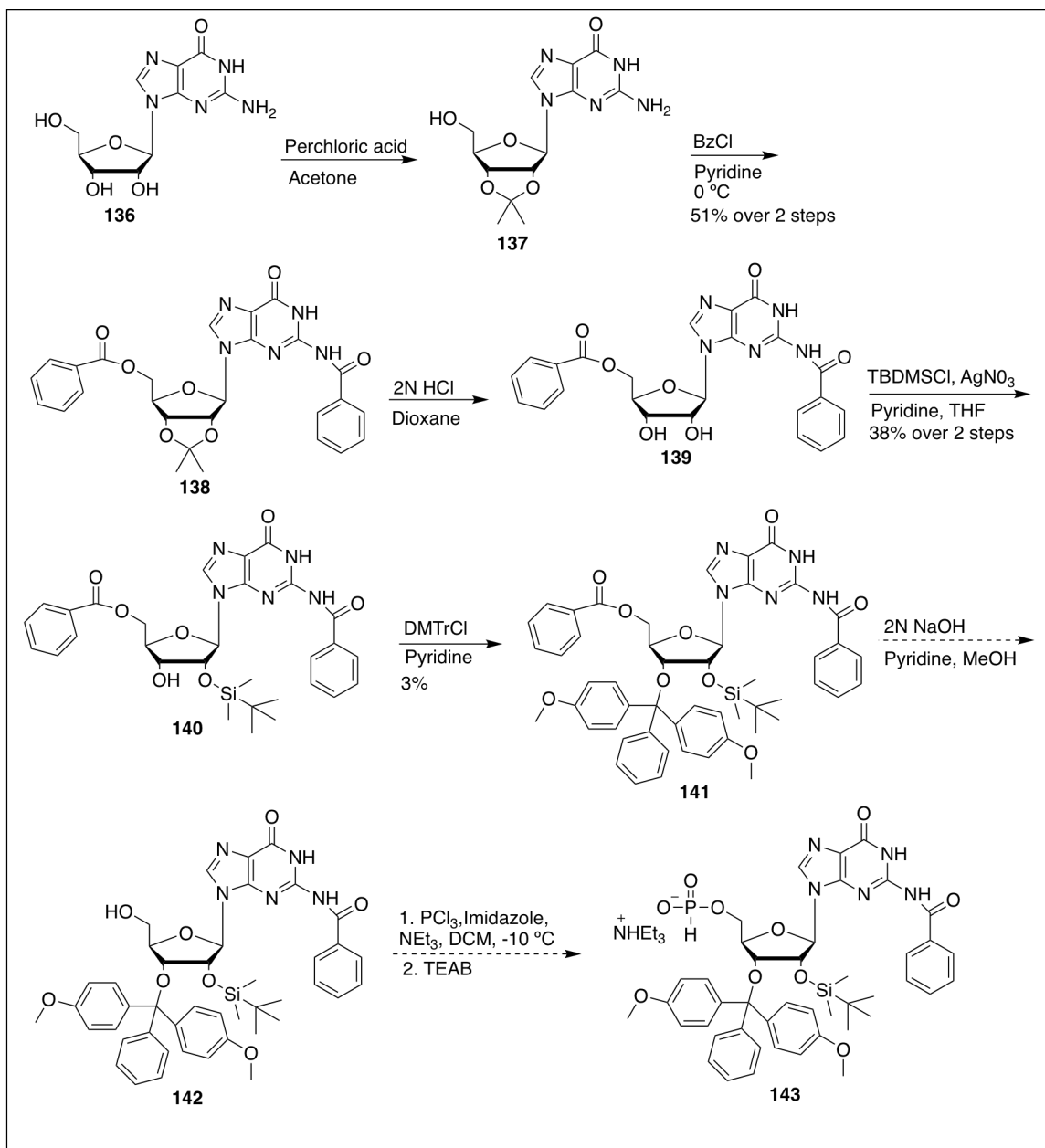
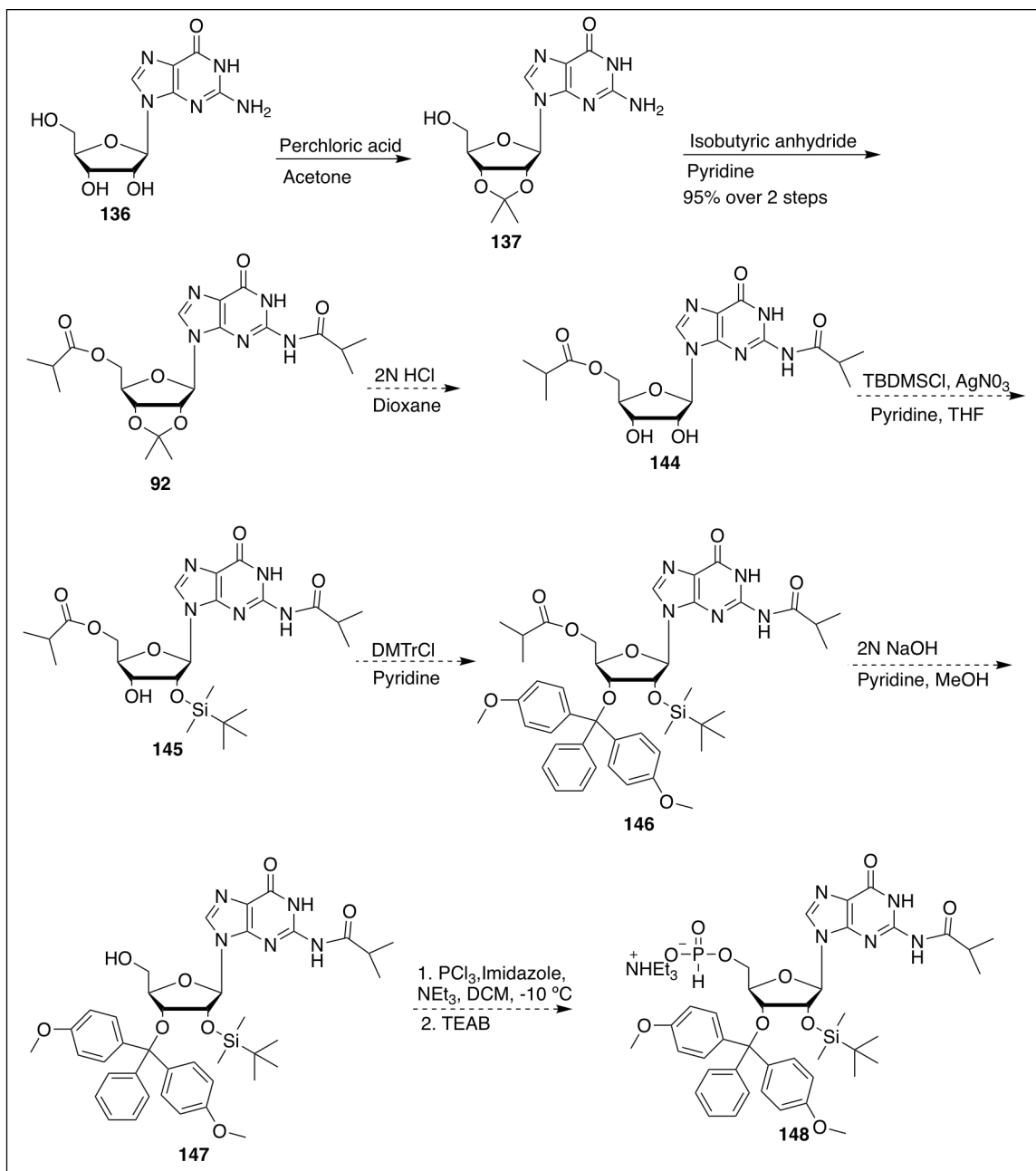


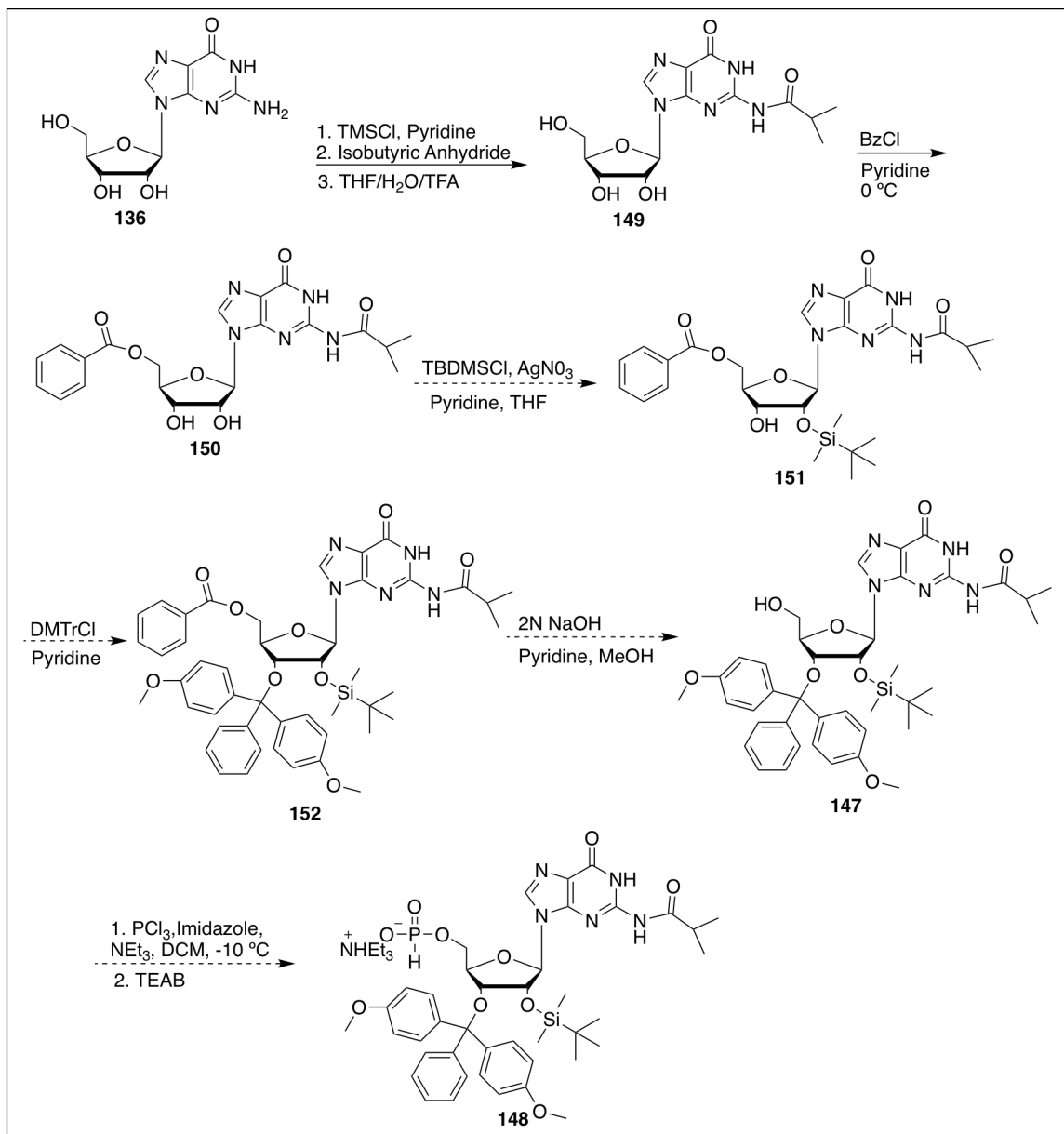
Figure 2-4. Guanosine Protecting Groups



Scheme 2-8. Attempted Synthesis of Guanosine 5' H-Phosphonate Using Benzoyl Protecting Groups



Scheme 2-9. Attempted Synthesis of Guanosine 5' H-Phosphonate Using Isobutyryl Protecting Groups



Scheme 2-10. Attempted Synthesis of Guanosine 5' H-Phosphonate Using a Combination of Isobutyryl and Benzoyl Protecting Groups

debenzoylation was attempted and appeared to work, however the compound degraded shortly after the purification.

Second, a similar approach as was used in the other nucleosides was attempted using the isobutyryl protecting groups, a more commonly used base protecting group for guanosine, seen in **Scheme 2-9**. Guanosine was protected using a catalytic amount of perchloric acid in acetone to give **137**¹⁰⁸. This was protected at the 5' position, as well as on the exocyclic amine, with Ibu using isobutyric anhydride in pyridine to afford **92**¹⁰⁹. Fully protected **92** underwent acidic hydrolysis using dilute hydrochloric acid in dioxane with the attempt of forming **139**¹⁰². However, under these conditions, the 5' isobutyryl group was consistently lost as well.

Finally, the approach outlined in **Scheme 2-10** was adopted. This removed the need to use the isopropylidene altogether, as well as utilized the more common Ibu protecting group on the base and the Bz group beneficial for solubility on the 5' position. The hydroxyl groups of guanosine were exposed to transient protection using TMSCl (trimethylsilyl chloride) in pyridine, followed by the addition of isobutyric anhydride to protect the base and then the addition of water and THF with TFA (trifluoroacetic acid) to secure the removal of the TMS groups to afford **149**^{105, 109}. In the future, adding methanol may be more beneficial than water yielding more methyl isobutyrate, which has a lower boiling point than isobutyric acid and may be easier to remove. This also may allow for the purification of this compound in the future, though it will certainly have solubility issues.

Chapter 3

Conclusions

The goal of this work was to study the reactivity of the C5'-uridylyl radical and the lesions formed from its generation. Herein, the synthesis of the C5'-uridylyl radical precursor **96** was greatly improved, increasing the yield of multiple reactions for the continued study of this compound. Using instrumentation such as HPLC and MS, the outcomes of the C5'-uridylyl radical were studied. Under anaerobic conditions it was shown that no UV-active reactive intermediates formed under physiological conditions and that the base release product uracil was the major product. Using aerobic conditions, it was found that the 5' aldehyde **105** was the predominant lesion formed, with some uracil and uridine, as well as a product we were unable to identify. Though pH played a role in the product distribution in the anaerobic studies, it did not appear to have an effect on the outcomes under aerobic conditions. A mechanism for the formation of **105** was proposed using a combination of the data from this work and other literature sources. The H-phosphonate C5'-uridylyl radical precursor was synthesized for incorporation into strands of RNA, however much of it degraded to form phosphate **94**. This work also began the journey towards a new automated RNA synthesis in the 5' to 3' direction with

the synthesis of uridine 5' H-phosphonate, as well as laying the groundwork for the future synthesis of cytidine, adenosine, and guanosine 5' H-phosphonates **127**, **135**, and **148**.

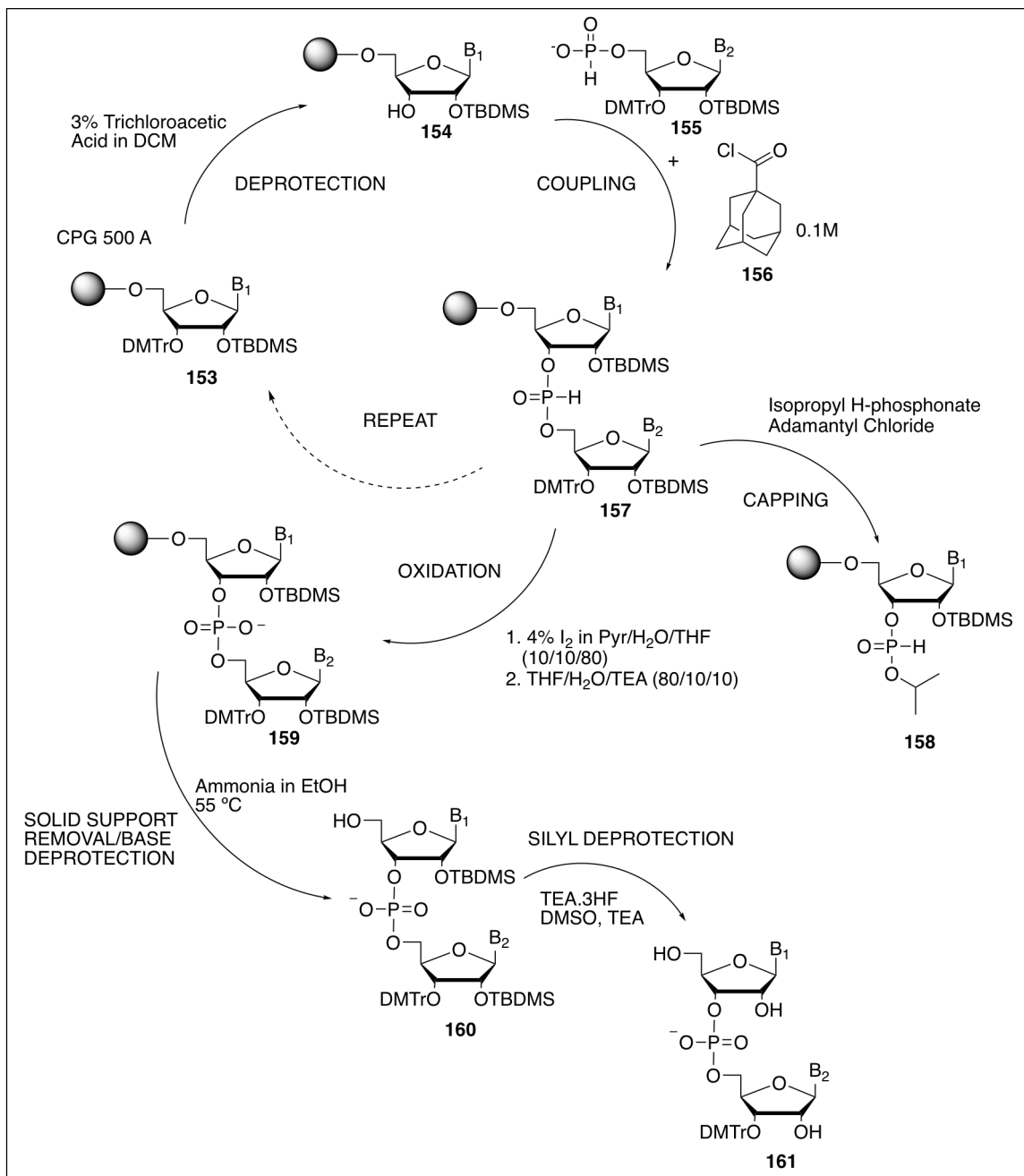
Chapter 4

Future Work

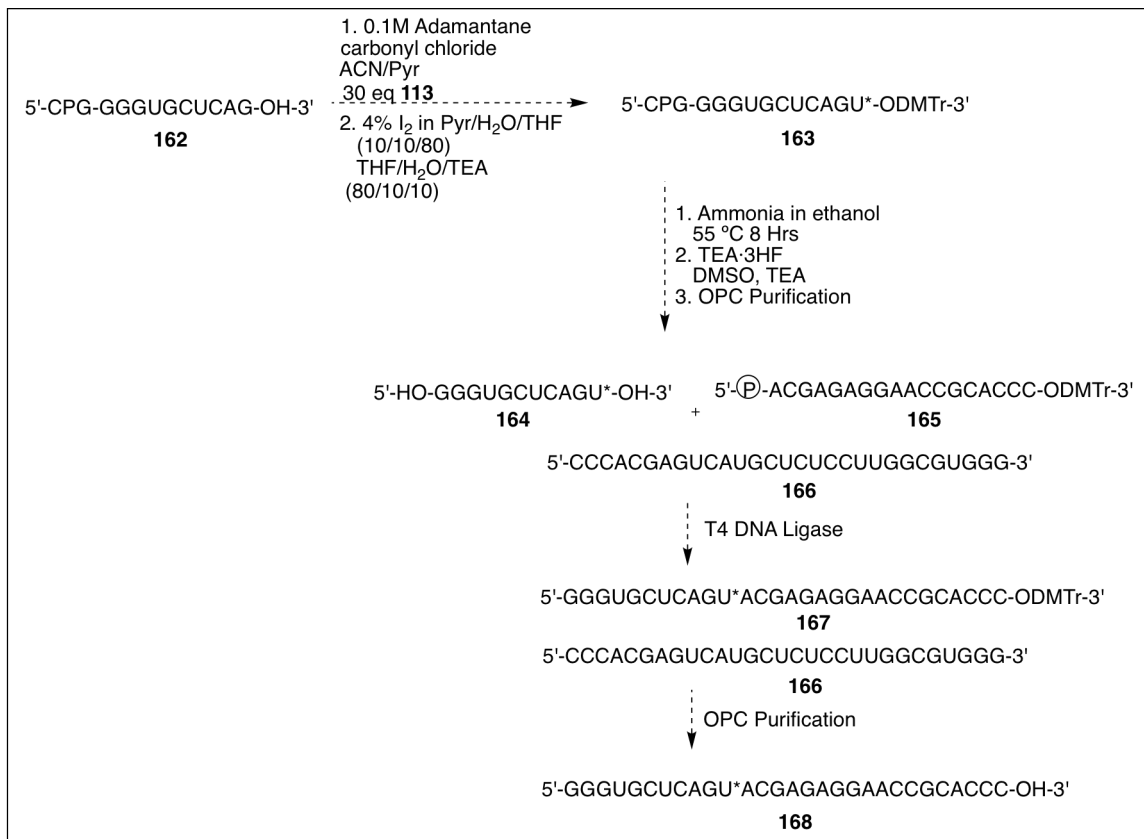
In order to further this work, the synthesis of cytidine, adenosine, and guanosine 5' H-phosphonates **127**, **135**, and **148** must be completed as described in Chapter 3. When all 4 H-phosphonates are in hand, a DNA synthesizer can be used to develop the automated RNA synthesis shown in **Scheme 4-1**. While the synthesis of these compounds are underway, the new automated RNA synthesis can begin development using the uridine 5' H-phosphonate that is in hand.

While the development of this automated synthesis is underway, more of H-phosphonate radical precursor **113** can be synthesized to incorporate into oligoribonucleotides, specifically, the Sarcin/Ricin Loop RNA. This RNA has already been used to study RNA oxidation and found that the 5' H atom was abstracted to form the C5' radical in U11³². Once obtained, hand-coupling techniques can be used to incorporate **113** into position 11 of a portion of the Sarcin/Ricin Loop. After purification, this can then be ligated with the remainder of the Sarcin/Ricin Loop RNA as seen in **Scheme 4-2**. Once isolated, the modified Sarcin/Ricin Loop RNA **168** can be subjected to photolysis conditions with excess GSH to prove that the radical is generated (**Scheme 4-3**). With excess GSH under anaerobic conditions, the radical should be reduced to give

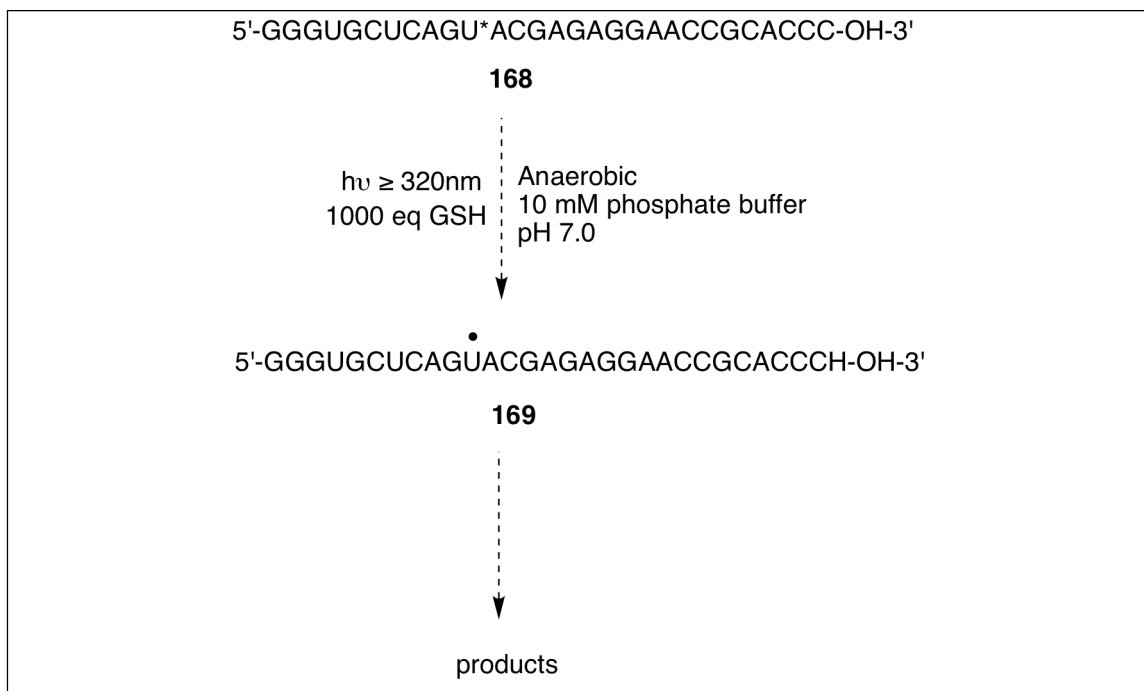
the unmodified Sarcin/Ricin Loop RNA. Once the method is tested, aerobic conditions can be used to compare results to Tullius' study³². When this proof of concept has been completed, **113** could be used in a number of strands of RNA to study the effect of RNA oxidation.



Scheme 4-1. Proposed Automated RNA Synthesis Approach



Scheme 4-2. Synthesis of Modified Sarcin/Ricin Loop RNA



Scheme 4-3. Photolysis of Modified Sarcin/Ricin Loop RNA

Chapter 5

Experimental Procedures

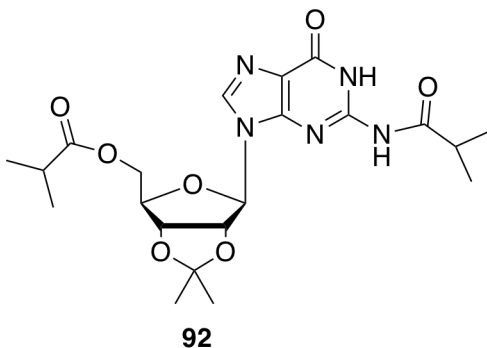
5.1 Materials, Methods, and Equipment

All reactions were carried out under laboratory conditions using argon or nitrogen in clean, oven-dried flask unless stated otherwise. All chemicals and reagents were obtained commercially and used without further purification unless otherwise stated. New compounds were characterized using Nuclear Magnetic Resonance (NMR) and High Resolution Mass Spectrometry (HRMS). ^1H NMR were obtained on a VXR 400, Inova 600, or Avance 600 NMR using CDCl_3 or MeOH-d_4 . ^{13}C NMR were obtained on an Avance 600 NMR using CDCl_3 or MeOH-d_4 . ^{31}P NMR were obtained on a VXR 400 using CDCl_3 . HRMS were obtained from Dr. Dragan Isalovic's laboratory using an electrospray ionization (ESI)-quadrupole time-of-flight (Q-TOF) mass spectrometer. Low Resolution Mass Spectrometry were obtained using either an Esquire-electrospray ionization mass spectrometer or a Shimadzu LCMS 2020. High-Performance Liquid Chromatography (HPLC) was completed using either a Dionex Ultimate 3000 equipped with a variable wavelength detector or a Shimadzu LC 20-AT with a UV detector. High-Performance Liquid Chromatography-Mass Spectrometry (LCMS) was accomplished

using a Shimadzu LC 20-AT connected to a Shimadzu LCMS 2020. Purification of compounds was done using either a manual column with silica or alumina or a Biotage SP4 automated chromatography system using Biotage SNAP pre-packaged silica columns. Thin Layer Chromatography (TLC) was carried out using silica gel 60 F254 aluminum backed plates and visualized by UV light at 254 nm followed by staining and burning using anisaldehyde dip. Photolysis reactions were carried out using a Thermo Oriel with a 500 W mercury lamp and cut-off filter for wavelengths greater or equal to 320 nm, a Thermo Haake K20 cooling system and a Varian SPV Cary single cell peltier accessories. Other equipment used include a solvent purification system (Innovative Technology PS-MD-2 Pure Solvent System), rotary evaporators (Heidolph Collegiate Brinkmann rotovap), high vacuum pump (Edwards RV3), speed vacuum (Thermo electron savant DNA120), micro centrifuge (Thermo Sorvall Legend Micro 21), vortex mixer (Fisher Scientific), Milli-Q water purification system, pH meter (Fisher Accumet Basic AB15), and pipettes (Eppendorf Series 2100).

5.2 Synthesis of Novel Compounds

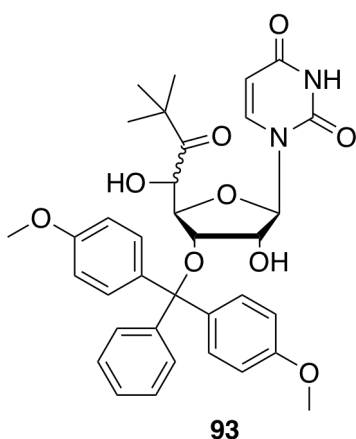
5.2.1 ((3*aR*,4*R*,6*R*,6*aR*)-6-(2-isobutyramido-6-oxo-1,6-dihydro-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro[3,4-*d*][1,3]dioxol-4-yl)methyl isobutyrate **92**



5.10 g (18.0 mmol) guanosine was dissolved in 245 mL of acetone and then 3.5 mL of perchloric acid was added dropwise while stirring. After 24 hours, the mixture was submerged in an ice bath and 3.0 mL of ammonium hydroxide was added dropwise. The media was filtered and the collected precipitate was washed with cold acetone to afford a white solid. The compound was coevaporated with toluene then dissolved in 210 mL of pyridine and then 25 mL of isobutyric anhydride were added slowly while stirring. After 25 hours, the reaction was quenched with 25 mL of water and concentrated. The compound was purified on a silica column using an eluent of 1% to 10% methanol (MeOH) in dichloromethane (DCM). 95% yield. ¹H NMR (600 Mz, CDCl₃) δ 12.12 (1H, s), 7.69 (1H, s), 5.99 (1H, d, J = 1.3), 5.30 (1H, dd, J = 11.1, 9.5), 5.07 (1H, dd, J = 6.2, 1.3), 4.96 (1H, dd, J = 6.5, 3.3), 4.44 (1H, m), 3.83, (1H, dd, J = 11.2, 5.8), 2.77 (1H, sep, J = 7.1), 2.65 (1H, sep, J = 7.1), 1.91 (1H, s), 1.57 (3H, s), 1.32 (3H, s), 1.29 (3H, d, J = 6.9), 1.26 (3H, d, J = 6.9), 1.21 (3H, d, J = 7.1), 1.19 (3H, d, J = 7.0). ¹³C NMR (600 Mz,

CDCl₃) δ 179.37, 178.62, 155.64, 148.14, 147.25, 138.43, 122.54, 114.47, 91.49, 86.21, 85.70, 82.27, 62.85, 36.56, 34.28, 27.28, 25.53, 19.36, 19.18, 19.05, 19.02. HRMS [M + H]⁺ calculated for C₂₁H₂₉N₅O₇ 464.2145. Found 464.2142.

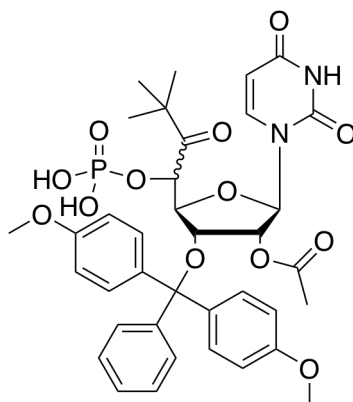
5.2.2 1-((2*R*,3*R*,4*S*,5*R*)-4-(bis(4-methoxyphenyl)(phenyl)methoxy)-3-hydroxy-5-(1-hydroxy-3,3-dimethyl-2-oxobutyl)tetrahydrofuran-2-yl)pyrimidine-2,4(1*H*,3*H*)-dione **93**



0.54 g (1.64 mmol) **96** was dissolved in 8 mL of pyridine then 0.65 g (1.92 mmol) dimethoxytrityl chloride (DMTrCl) was added. The reaction was stirred at room temperature. After 74 hours, an additional 0.10 g (0.30 mmol) dimethoxytrityl chloride and 0.02 g (0.16 mmol) DMAP (4-dimethylaminopyridine) were added to the solution. The reaction was chilled in an ice bath and quenched with 0.2 mL of methanol after 115 hours then concentrated. The compound was purified on a silica column using an eluent of 50% to 60% ethyl acetate (EA) in hexane with 5% triethylamine (NEt₃). 57% yield. ¹H NMR (600 Mz, MeOH-d₄) δ 7.65 (1H, d, J = 8.0), 7.63 (1H, d, J = 8.0), 7.47 (2H, d, J = 8.0), 7.47 (2H, d, J = 8.0), 7.37 (2H, d, J = 8.9), 7.37 (2H, d, J = 8.5), 7.33 (2H, d, J =

8.9), 7.32 (2H, d, J = 8.9), 7.21 (3H, t, J = 7.5), 7.20 (3H, t, J = 8.5), 6.82 (2H, d, J = 9.1), 6.80 (2H, d, J = 8.8), 6.77 (2H, d, J = 8.1), 6.76 (2H, d, J = 8.8), 6.31 (1H, d, J = 7.8), 6.27 (1H, d, J = 7.8), 5.49 (1H, d, J = 8.2), 5.48 (1H, d, J = 8.2), 4.59 (1H, d, J = 2.9), 4.52 (1H, d, J = 2.2), 4.46 (1H, dd, J = 7.9, 4.6), 4.45 (1H, d, J = 2.1), 4.34 (1H, dd, J = 7.5, 4.7), 4.27 (1H, d, J = 2.5), 3.77 (3H, s), 3.74 (3H, s), 3.74 (3H, s), 3.72 (3H, s), 3.23 (1H, d, J = 4.7), 2.86 (1H, d, J = 4.7), 1.12 (9H, s), 1.11 (9H, s). ¹³C NMR (600 Mz, MeOH-d₄) δ 214.98, 214.75, 166.37, 160.59, 160.53, 160.48, 160.42, 152.97, 152.84, 146.77, 146.72, 142.97, 142.78, 137.05, 136.05, 136.84, 136.79, 131.94, 131.82, 131.64, 131.60, 129.39, 129.34, 129.02, 128.95, 128.09, 126.43, 114.51, 114.48, 114.33, 103.68, 103.52, 88.80, 88.77, 88.26, 88.04, 87.50, 87.40, 77.90, 77.33, 75.15, 74.87, 72.99, 70.54, 55.80, 44.50, 44.10, 27.34, 26.81. HRMS [M + Na]⁺ calculated for C₃₅H₃₈N₂O₉ 653.2475. Found 653.2484.

5.2.3 (2*R*,3*R*,4*R*,5*S*)-4-(bis(4-methoxyphenyl)(phenyl)methoxy)-5-(3,3-dimethyl-2-oxo-1-(phosphonoxy)butyl)-2-(2,4-dioxo-3,4-dihydropyrimidin-1(2*H*)-yl)tetrahydrofuran-3-yl acetate **94**

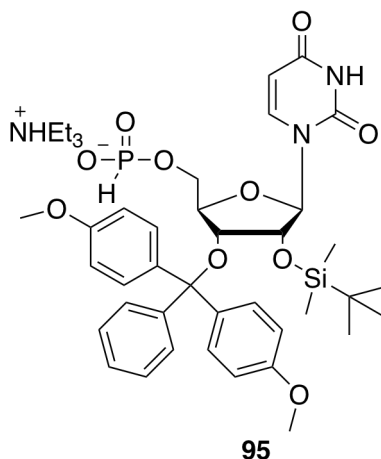


94

0.06 g (0.095 mmol) **93** and 2.50 mg (0.02 mmol) DMAP were dissolved in 8.0 mL of DCM. 10 μ L of acetic anhydride and 20 μ L of triethylamine were added and the reaction was stirred for 126 hr until quenched with methanol and concentrated. The reaction mixture was run through a silica column using an eluent of 30% to 40% ethyl acetate in hexane with 5% triethylamine then 1% to 3% methanol in 40/60 ethyl acetate/hexane with 5% triethylamine. After concentrating, the crude product was used directly in the next step. 0.07 g (1.03 mmol) imidazole was coevaporated with acetonitrile twice then dissolved in 3.0 mL of DCM and submerged in an ice salt bath. 0.03 mL of phosphorous trichloride (PCl_3) was added slowly followed by 0.15 mL of triethylamine in 0.15 mL of DCM. This solution was allowed to stir for 30 min. Then 0.06 g (0.089 mmol) of the nucleoside was coevaporated separately in toluene twice and dissolved in 2.0 mL of DCM. This was added dropwise to the stirring solution. After 1 hour, the flask was removed from the ice bath and 6.0 mL of triethylammonium bicarbonate buffer (TEAB)

was added. The organic layer was separated and washed with an equal volume of triethylammonium bicarbonate buffer again then concentrated. The H-phosphonate compound was purified on a silica column using an eluent of 0% to 8% methanol in dichloromethane with 5% triethylamine, leading to its oxidation. 27% yield. ^1H NMR (600 Mz, CDCl_3) δ 7.40 (1H, d, $J = 1.9$), 7.39 (1H, d, $J = 1.4$), 7.35 (1H, d, $J = 8.3$), 7.30 (2H, d, $J = 9.1$), 7.29 (2H, d, $J = 8.6$), 7.24 (3H, t, $J = 5.7$), 6.77 (2H, d, $J = 3.5$), 6.75 (2H, d, $J = 3.5$), 6.46 (1H, d, $J = 8.3$), 6.27 (1H, d, $J = 8.4$), 5.66 (1H, d, $J = 8.3$), 5.59 (1H, d, $J = 8.3$), 5.46 (1H, d, $J = 3.8$), 5.43 (1H, d, $J = 2.4$), 4.38 (1H, d, $J = 5.6$), 4.36 (1H, d, $J = 2.2$), 4.23 (1H, dd, $J = 8.2, 5.2$), 3.77 (3H, s), 3.76 (3H, s), 3.63 (6H, q, $J = 7.5$), 3.59 (1H, d, $J = 5.5$), 2.17 (3H, s), 2.10 (3H, s), 2.04 (3H, s), 1.99 (3H, s), 1.47 (9H, t, $J = 7.0$), 1.13 (9H, s), 1.12 (9H, s). ^{13}C NMR (600 Mz, CDCl_3) δ 207.21, 206.66, 170.63, 168.51, 162.37, 159.12, 158.75, 150.46, 144.11, 139.57, 139.10, 135.15, 135.06, 130.10, 130.07, 129.26, 128.13, 128.12, 127.99, 127.89, 127.63, 127.22, 113.41, 113.39, 113.29, 103.62, 87.38, 86.03, 81.51, 74.73, 74.41, 72.93, 55.44, 55.40, 53.13, 45.97, 43.53, 26.83, 26.48, 21.11, 20.65, 8.28. ^{31}P NMR (400 Mz, CDCl_3) δ 47.9, 9.3. HRMS $[\text{M} + \text{H}]^+$ calculated for $\text{C}_{37}\text{H}_{41}\text{N}_2\text{O}_{12}\text{P}$ 737.2475. Found 737.1410.

5.2.4 ((2*R*,3*R*,4*R*,5*R*)-3-(bis(4-methoxyphenyl)(phenyl)methoxy)-4-((*tert*-butyldimethylsilyl)oxy)-5-(2,4-dioxo-3,4-dihydropyrimidin-1(2*H*)-yl)tetrahydrofuran-2-yl)methyl phosphonate triethylammonium salt **95**



0.28 g (5.88 mmol) imidazole was coevaporated in toluene three times then dissolved in 10.0 mL of DCM and submerged in an ice salt bath. 0.15 mL of phosphorous trichloride was added dropwise followed by 0.85 mL of triethylamine. The reaction was left to stir for 30 min then 0.21 g (0.50 mmol) 2'-*O*-TBDMS-3'-*O*-DMTr-Uridine was coevaporated separately three times in toluene and dissolved in 7.0 mL of DCM before adding to the solution dropwise. Reaction was left stirring for 85 min and then brought to room temperature and quenched with 25 mL of TEAB. The DCM layer was separated and washed with an equal volume of TEAB again then concentrated. Compound was purified on a silica column using an eluent of 0 – 8% methanol in dichloromethane with 3% triethylamine. 99% yield. ¹H NMR (400 Mz, CDCl₃) δ 8.01 (1H, d, J = 8.1), 7.54 (2H, d, J = 7.5), 7.41 (4H, d, J = 8.8), 7.25 (3H, t, J = 8.1), 6.79 (4H, d, J = 8.1), 6.29 (1H, d, J =

8.1), 5.69 (1H, d, J = 8.1), 4.15 (1H, dd, J = 6.4, 5.0), 4.04 (1H, d, J = 5.0), 3.76 (6H, s), 3.74 (1H, d, J = 4.9), 3.56 (1H, d, J = 7.1), 3.38 (1H, unresolved d), 2.90 (6H, q, J = 14.1, 7.1), 1.23 (9H, t, J = 7.3), 0.92 (9H, s), 0.05 (3H, s), -0.06 (3H, s). ¹³C NMR (600 Mz, CDCl₃) δ 163.29, 158.83, 150.75, 145.67, 141.50, 136.98, 136.56, 113.40, 102.62, 87.85, 86.72, 81.97, 74.90, 74.31, 63.15, 55.43, 45.76, 29.89, 26.05, 9.48, -4.17, -4.38. ³¹P NMR (400 Mz, CDCl₃) δ 5.6. HRMS [M + H]⁺ calculated for C₄₂H₆₀N₃O₁₀Psi 826.3858. Found 826.3860.

5.3 Photolysis of 96

5.3.1 Anaerobic Photolysis

Photolyses were carried out in plastic cuvettes (1 cm path length) with a 500 W mercury lamp with IR and cut-off filters and a focusing lens (≥ 320 nm) under argon. Reaction mixtures of 200 μ L contained 37 nmol of **96** in 10 mM phosphate buffer at pH 7.0 and 8 mM GSH (glutathione) and were photolyzed at room temperature for varying amounts of time (1, 2, 4, 8, 15, 30, and 60 min) after first bubbling argon through the mixture for 15 min. After photolysis, mixtures were injected directly into a Dionex Ultimate 3000 HPLC system using a Hypersil BDS C18 RP column (5 μ m, 4.6 x 250 cm). Solvent A was 50 mM TEAA (triethylammonium acetate) buffer at pH 7.0 and solvent B was acetonitrile. The program used was as follows: 0.2% B for 7 min, 0.2–22% B over 5 min, 22–0.2% B over 8 min. The flow rate was 1.0 mL/min and the temperature was 35 °C. Wavelength detection was at 254 and 260 nm. Compound identity was

confirmed using independent standards, as were the RP-HPLC (reverse phase-HPLC) calibration curves.

5.3.2 Aerobic Photolysis

Experiments were performed in plastic cuvettes (1 cm path length) on 50 nmol **96** in 10 mM ammonium acetate buffer at pH 7.0 or 3.5 and 8 mM GSH using a 500 W mercury lamp (≥ 320 nm) with IR and cut-off filters and a focusing lens under oxygen. All samples were subjected to photolysis for 30 min after bubbling oxygen through the solution for 15 min. Samples were analyzed on a Shimadzu LCMS 2020 using RP-HPLC and ESI MS. A Hypersil BDS C18 column (5 μ m, 4.6 x 250 mm) was used. Solvent A was water with 1% acetic acid and solvent B was acetonitrile with 0.1% acetic acid. The program used was as follows: 0-4% B over 25 mins, 4-98% B over 10 mins, 98% B for 10 mins, 98-0% B over 5 mins, 0% B for 10 mins. The flow rate was 0.3 mL/min, the column oven temperature was 30 °C, and the wavelength detector was at 254 nm. For the ESI MS, the heat block was 400 °C and drying gas was at 15 L/min. Data was collected in both positive and negative ion modes between 50 and 2000 m/z. Compound identification was determined using independent standards, as were the RP-HPLC calibration curves.

Appendix

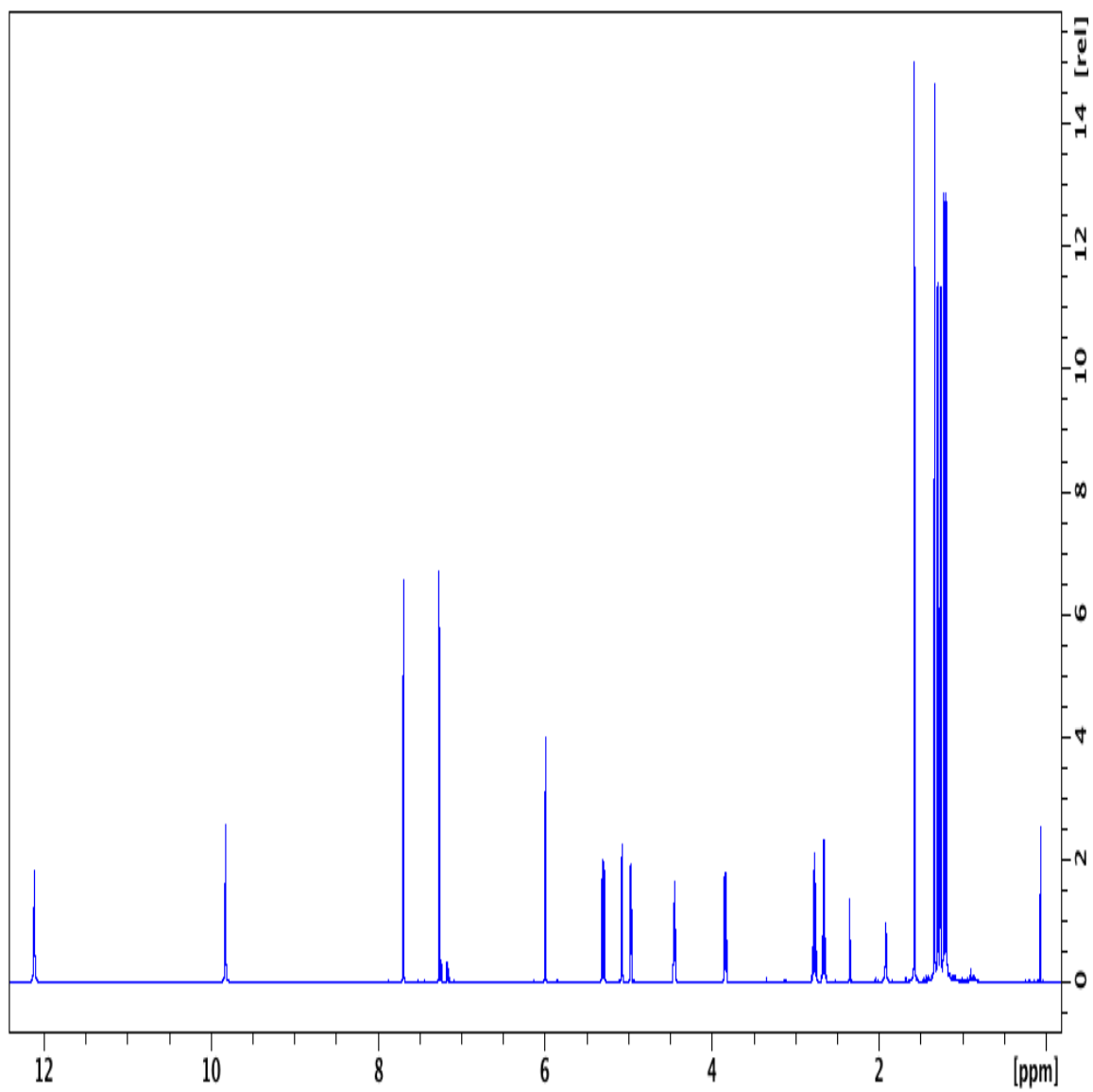


Figure A-1. ^1H NMR for **92**

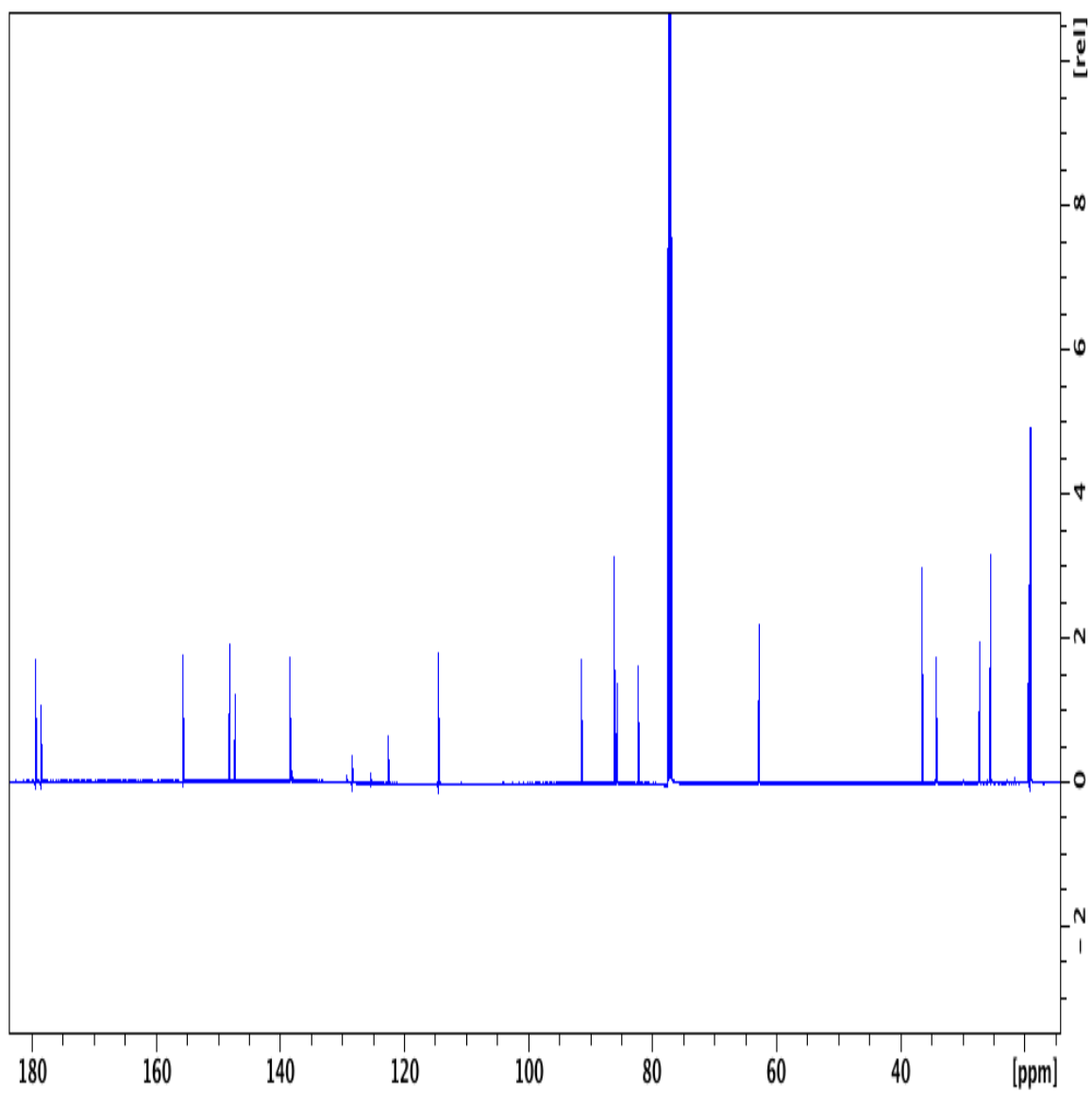


Figure A-2. ^{13}C NMR for **92**

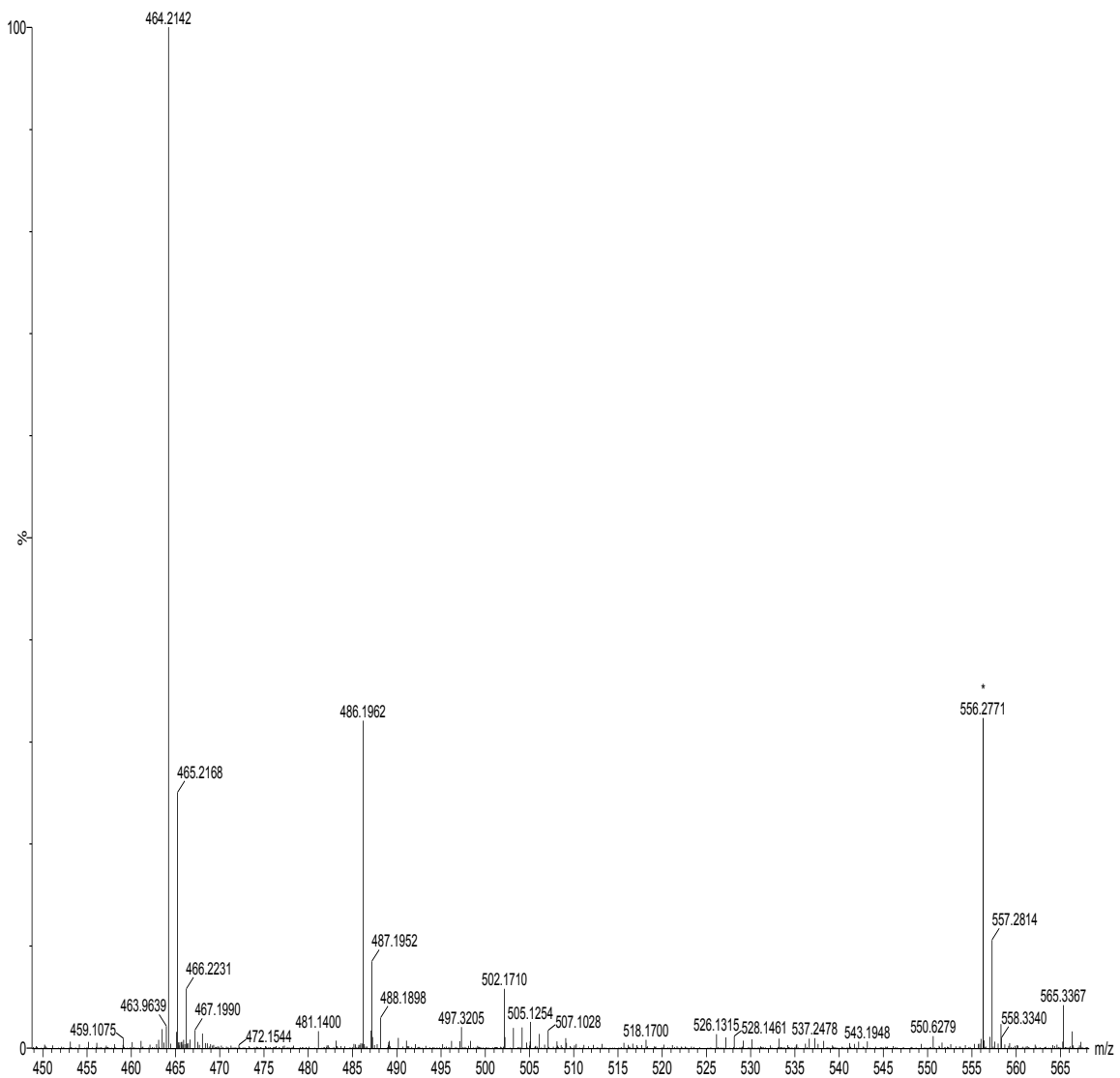


Figure A-3. HRMS for **92**

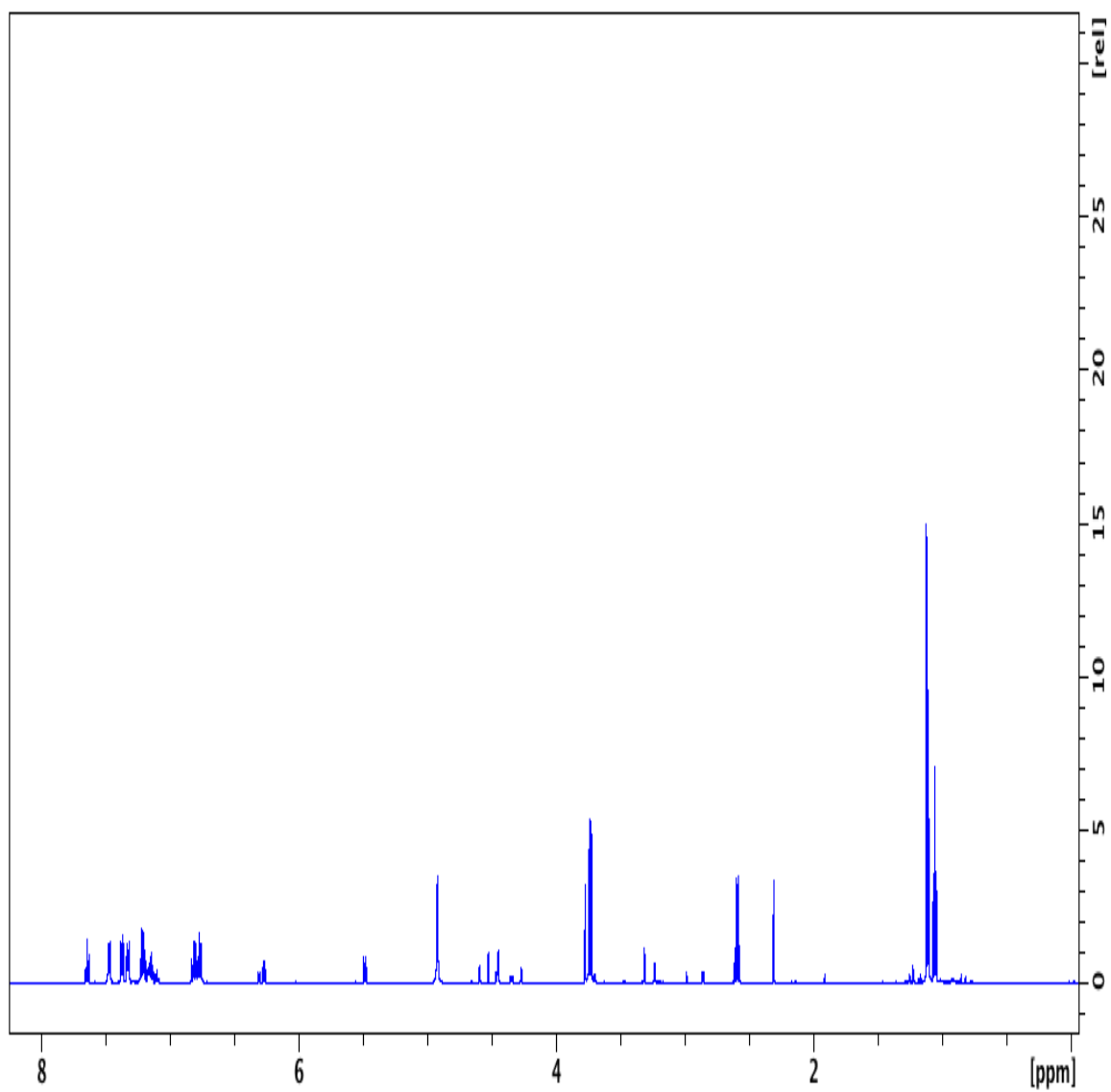


Figure A-4. ^1H NMR for **93**

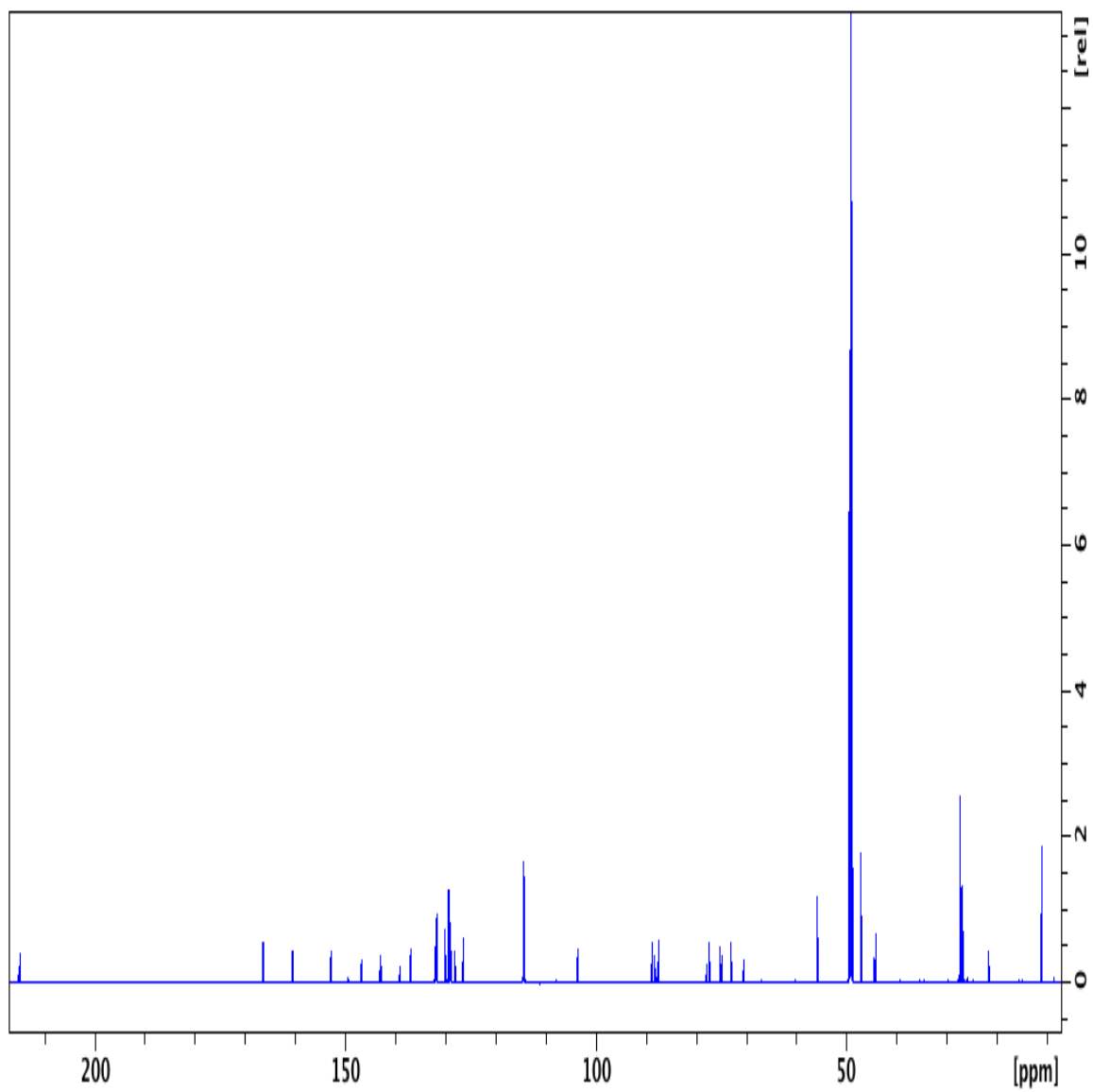


Figure A-5. ^{13}C NMR for **93**

KR_ME014_02222016 8 (0.153) AM (Cen,4, 80.00, Ar,8000.0,556.28,0.70); Cm (2:9)

TOF MS ES+
1.24e4

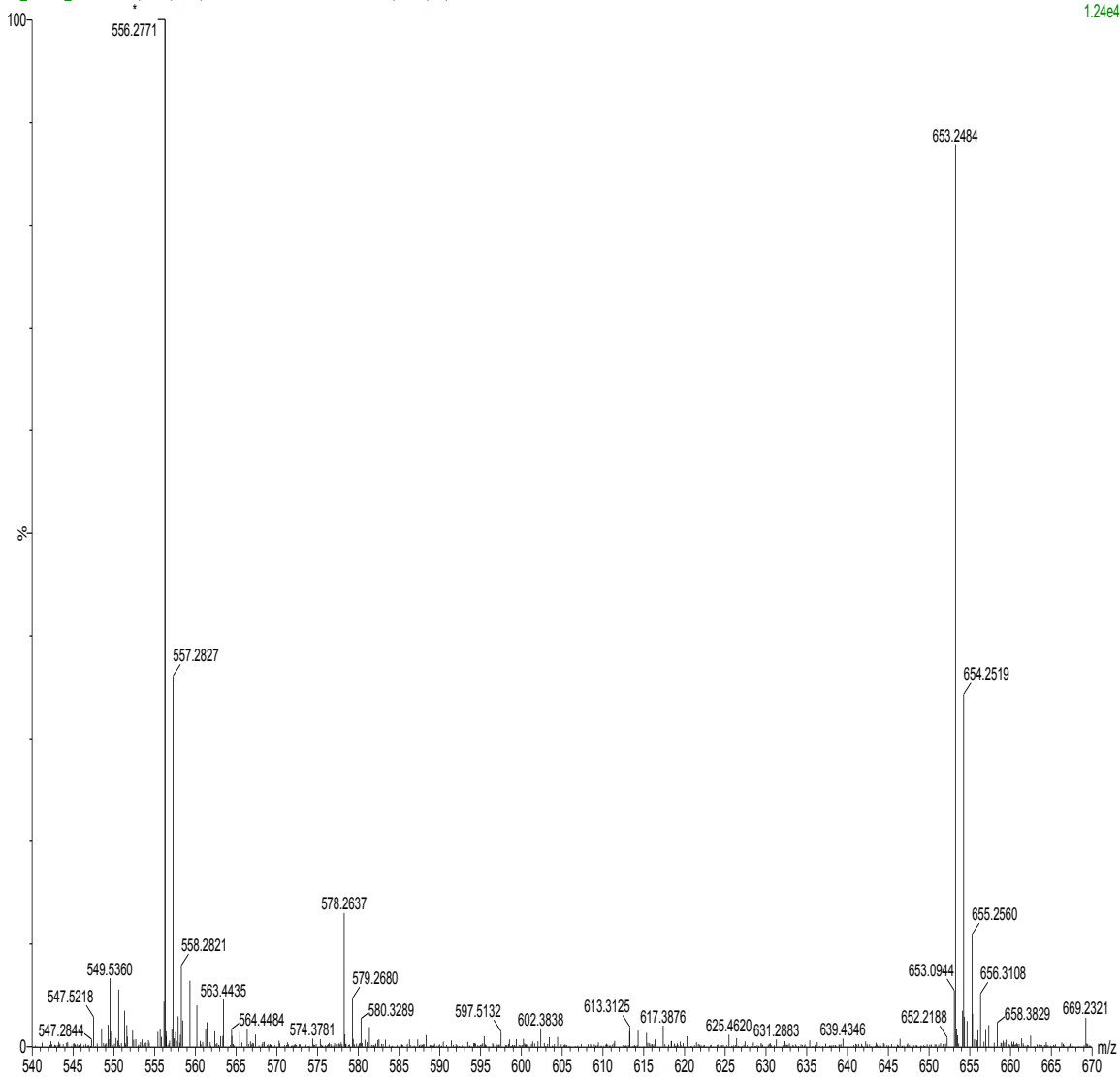


Figure A-6. HRMS for **93**

ME017 precursor phosphate

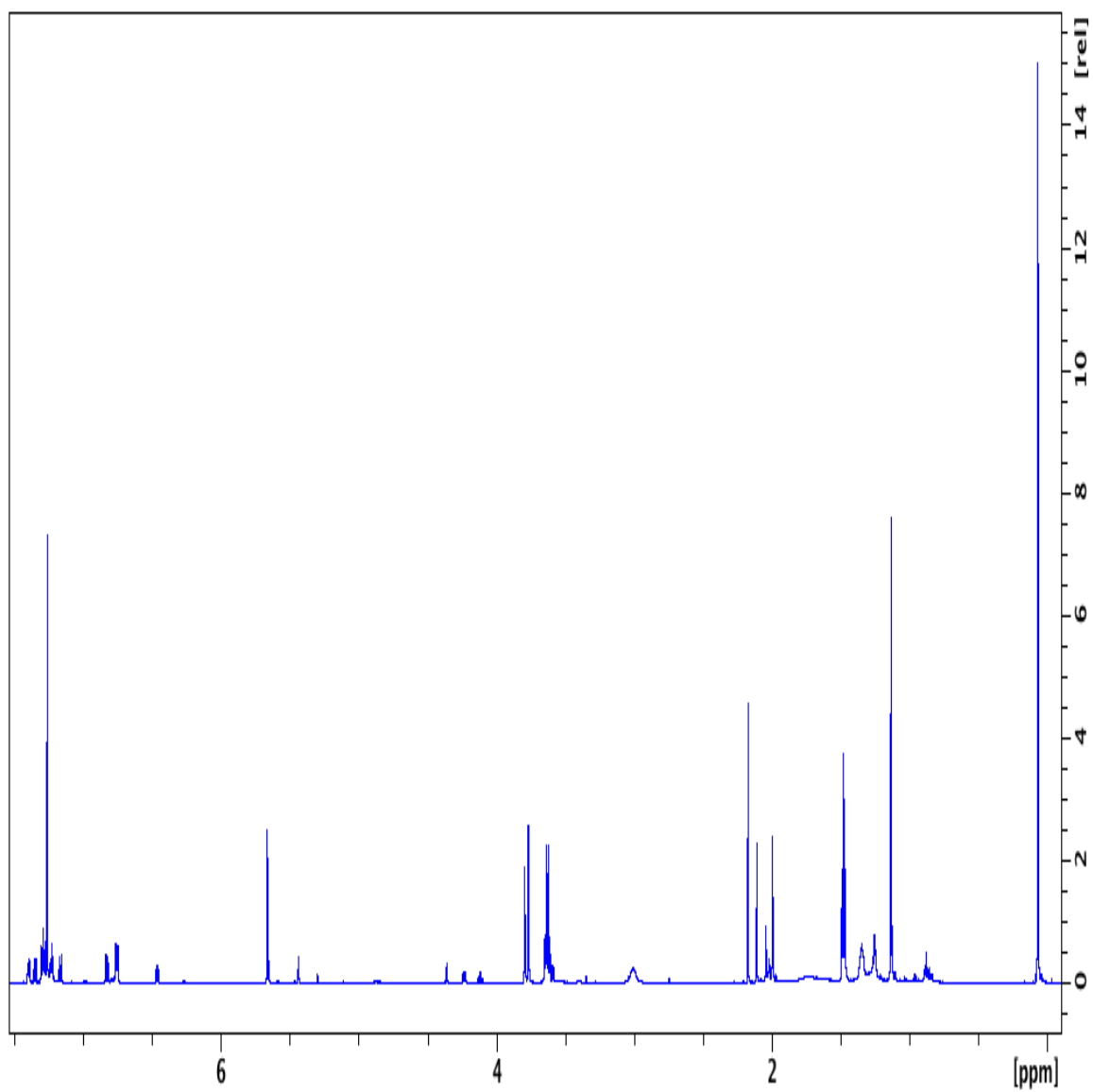


Figure A-7. ^1H NMR for **94**

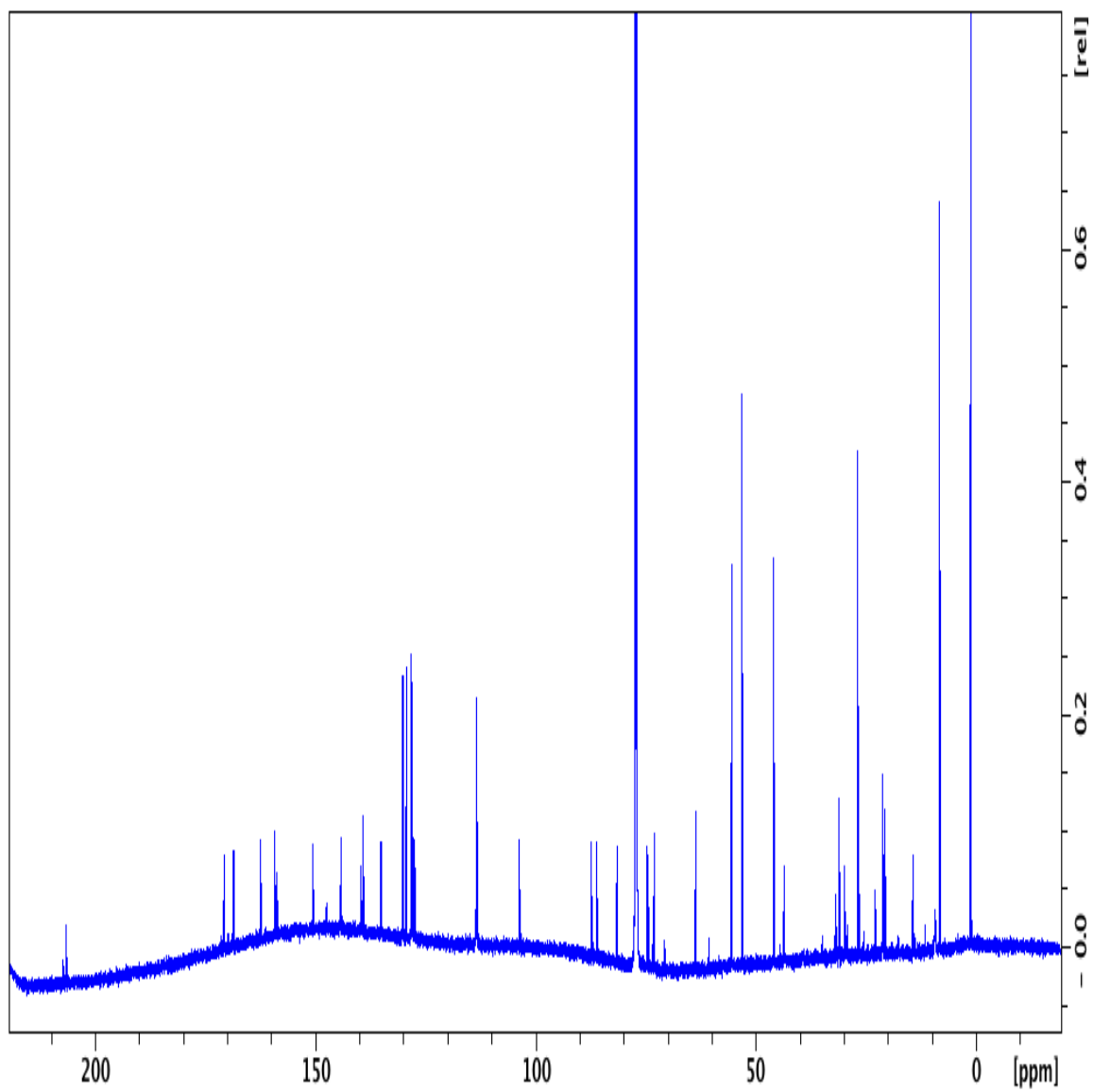


Figure A-8. ^{13}C NMR for **94**

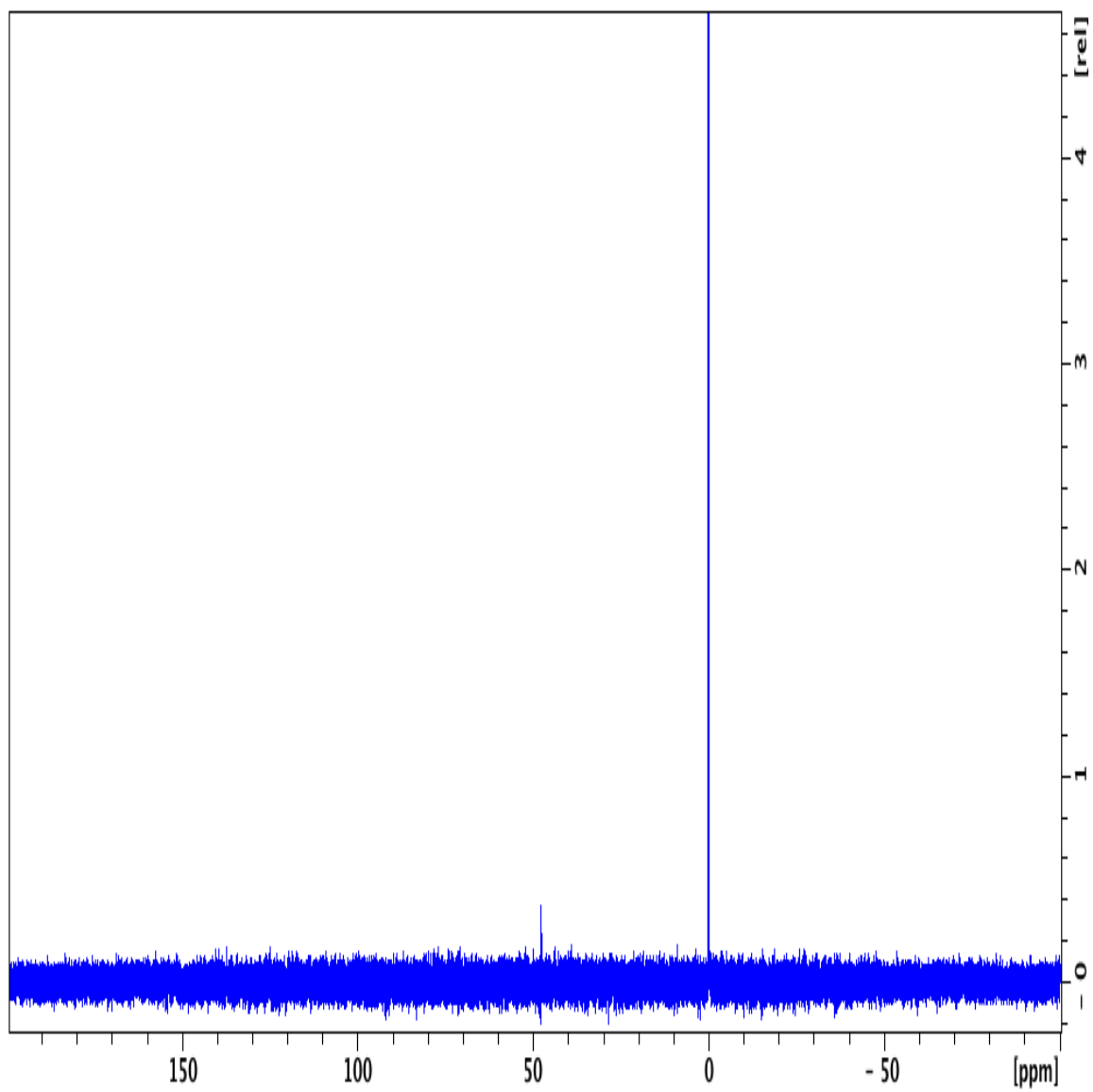


Figure A-9. ^{31}P NMR for **94**

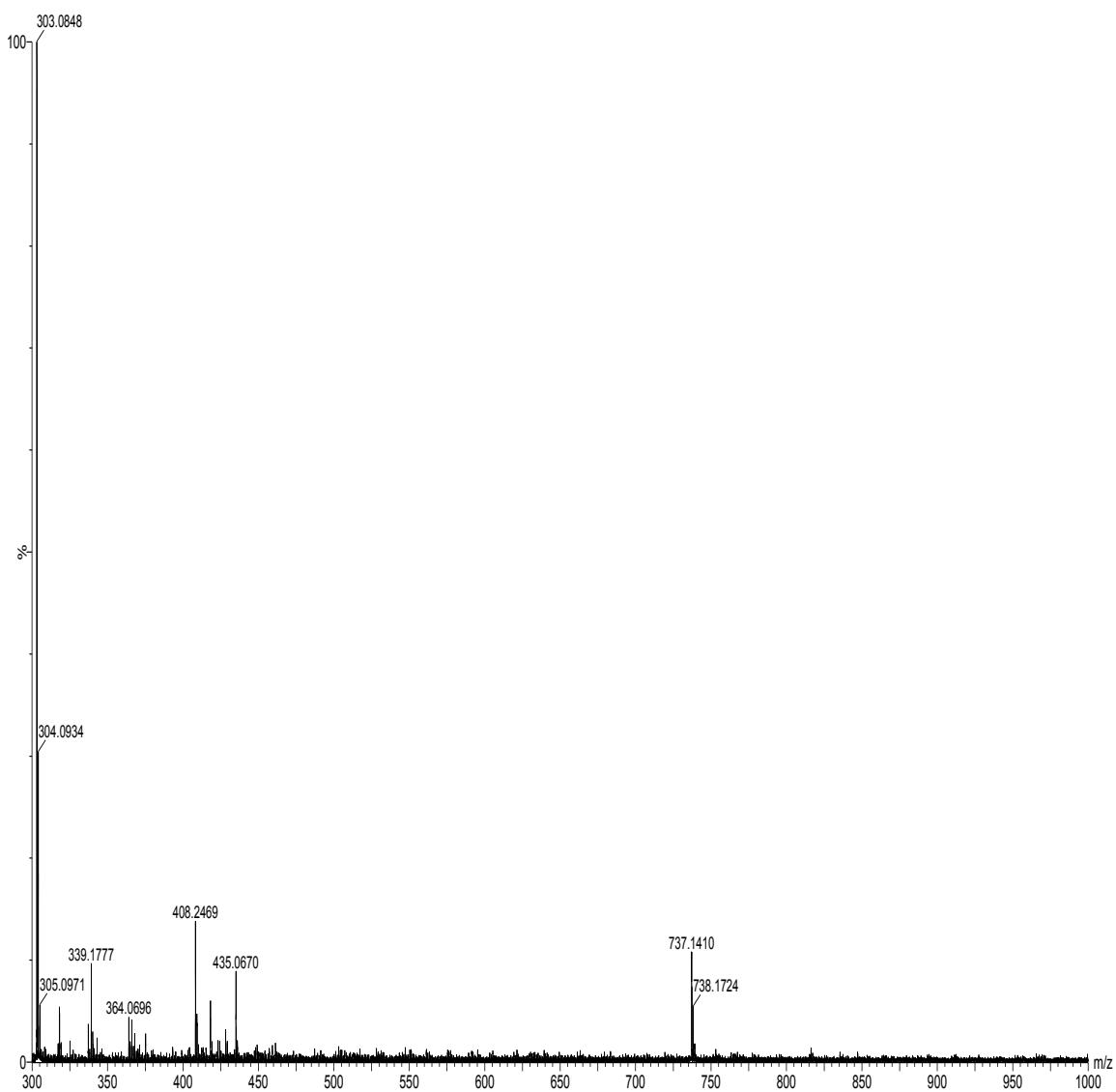


Figure A-10. HRMS for **94**

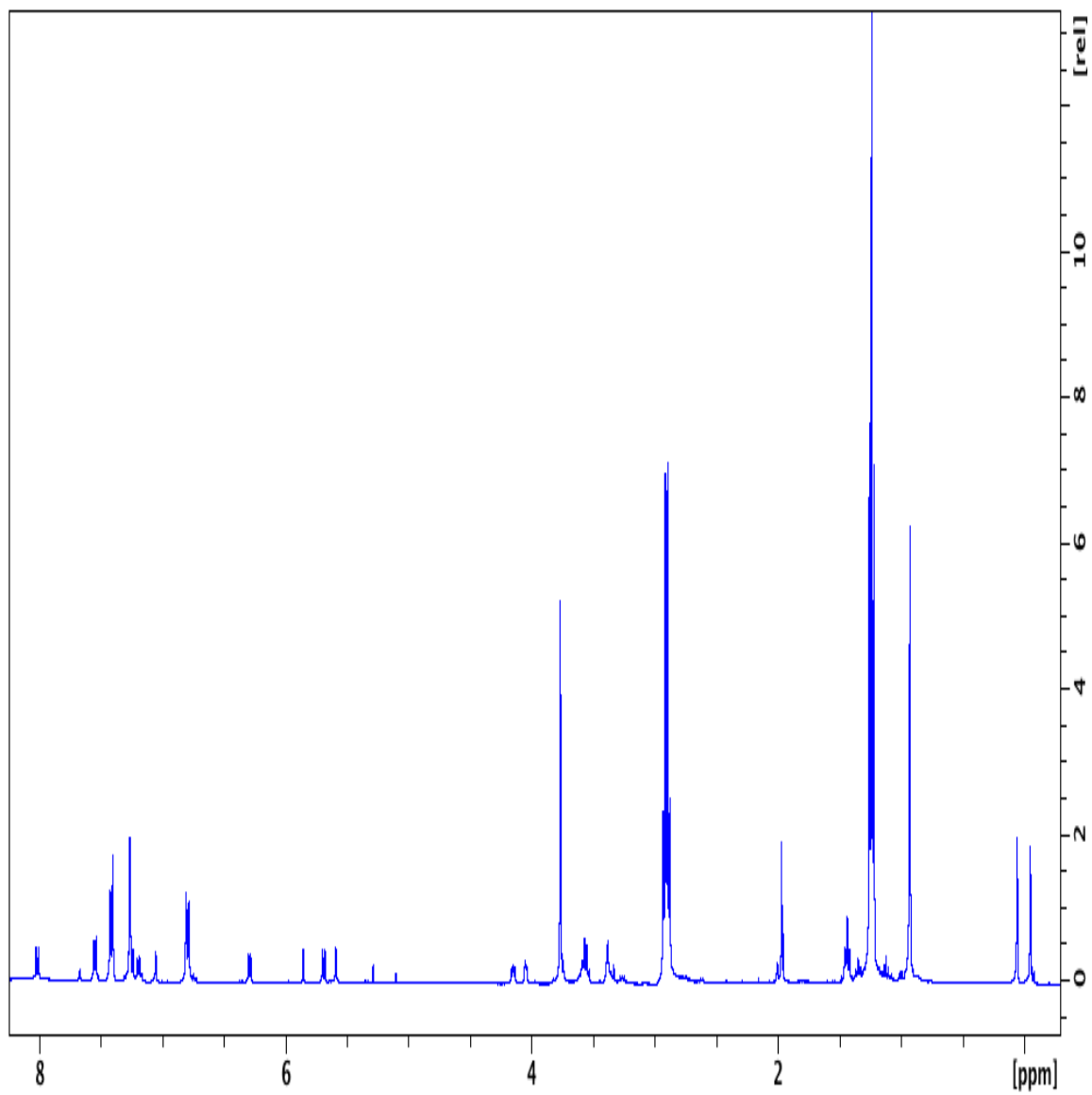


Figure A-11. ^1H NMR for 95

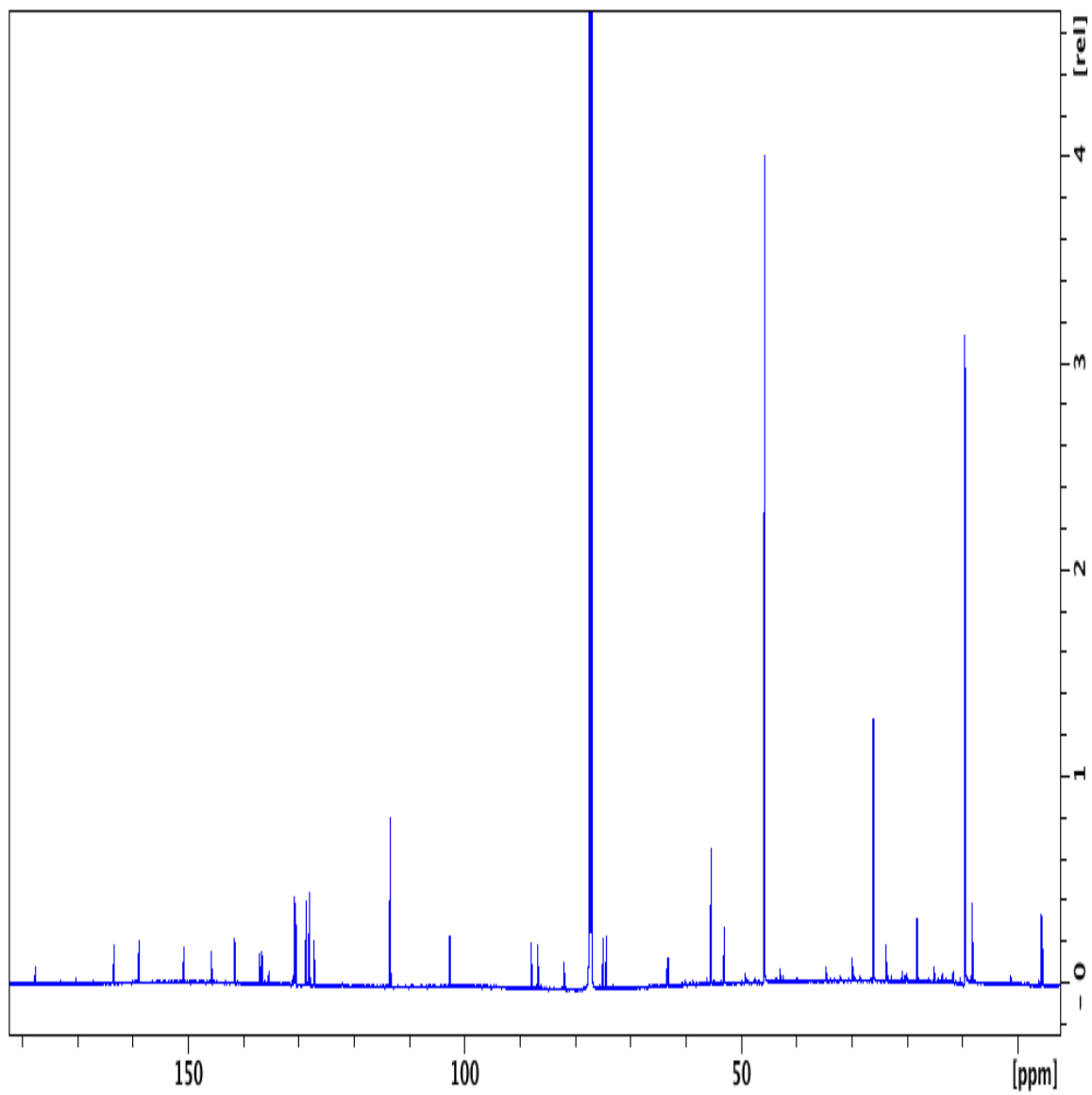


Figure A-12. ^{13}C NMR for **95**

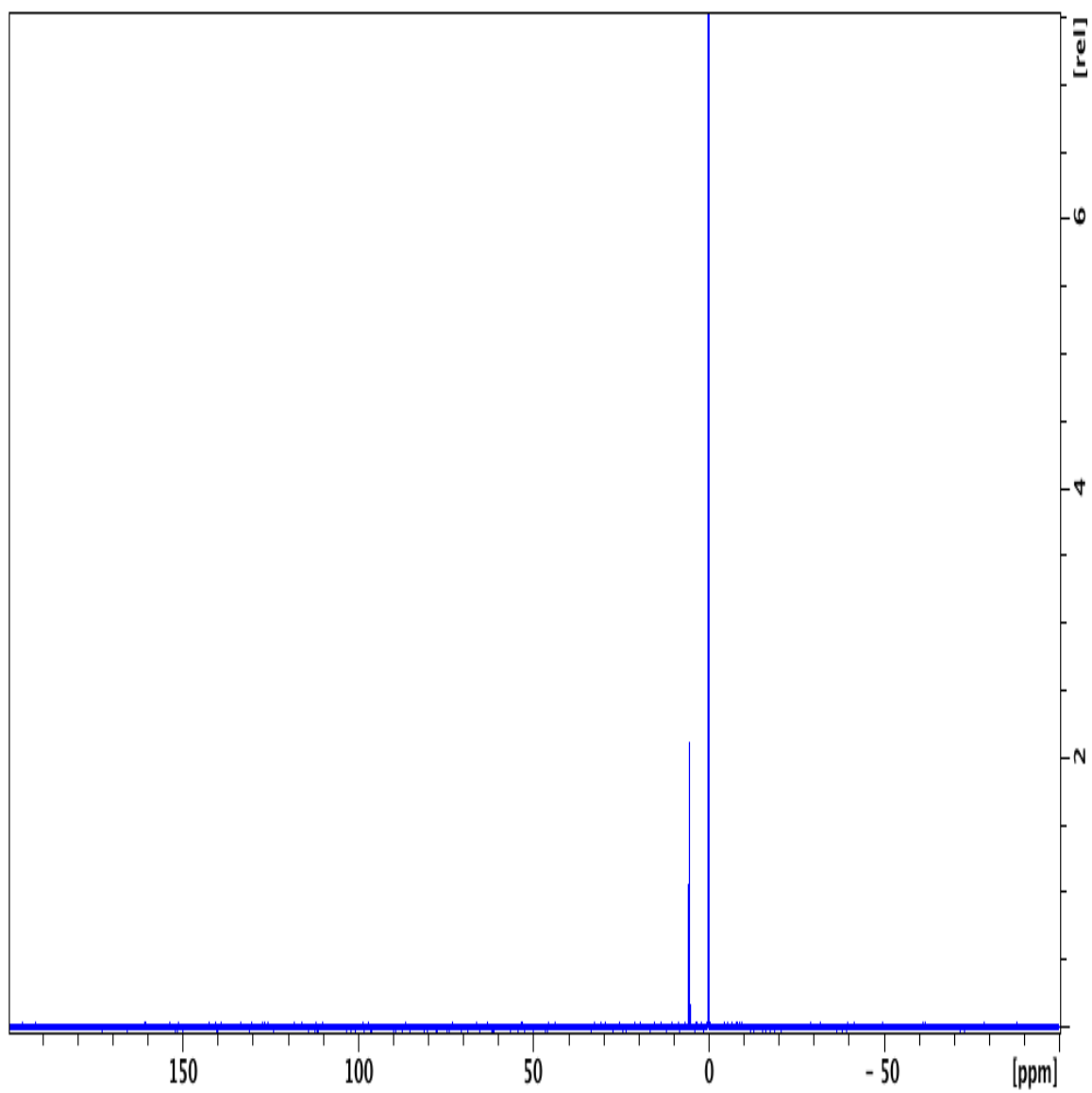


Figure A-13. ^{31}P NMR for **95**

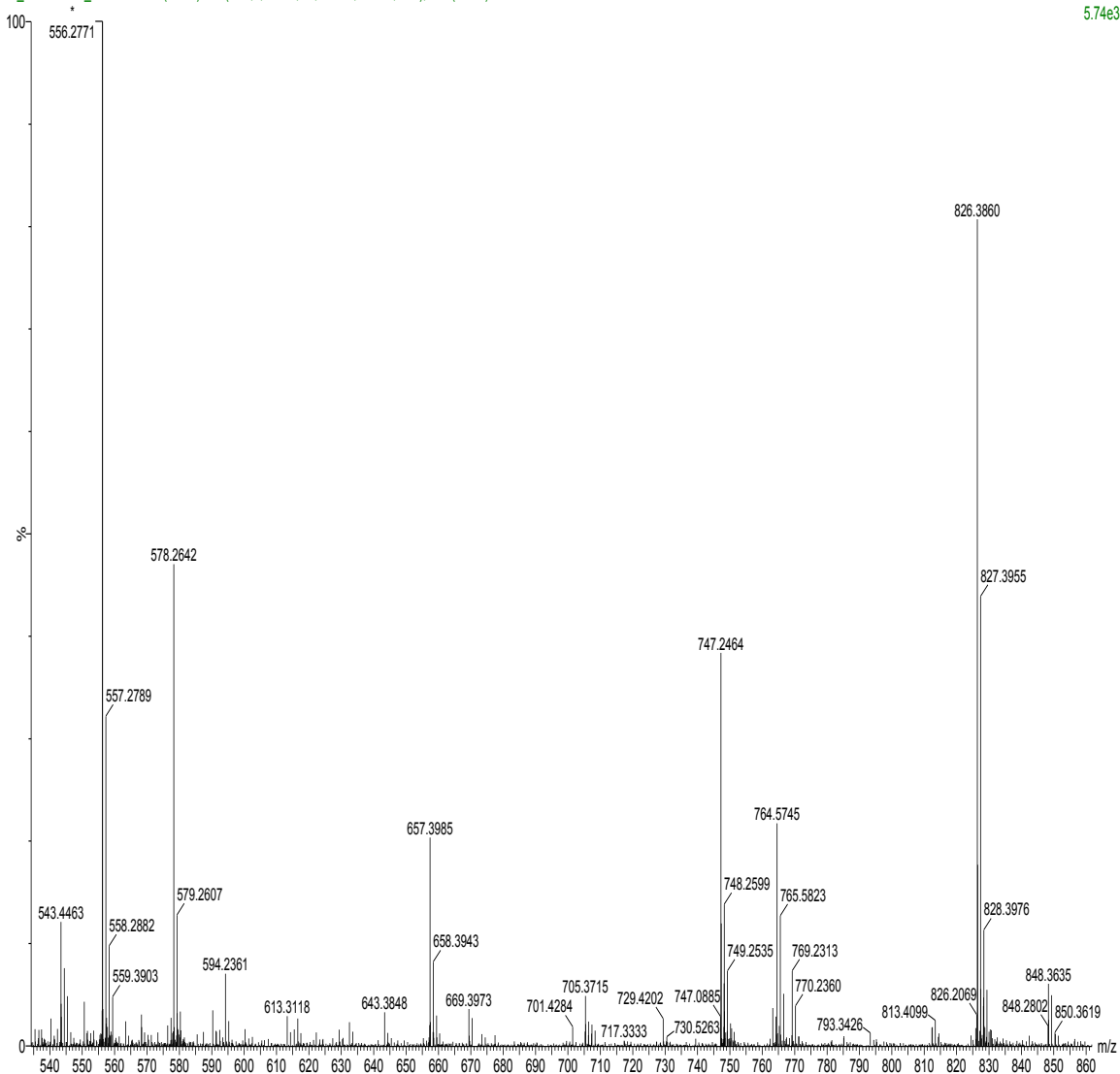


Figure A-14. HRMS for **95**

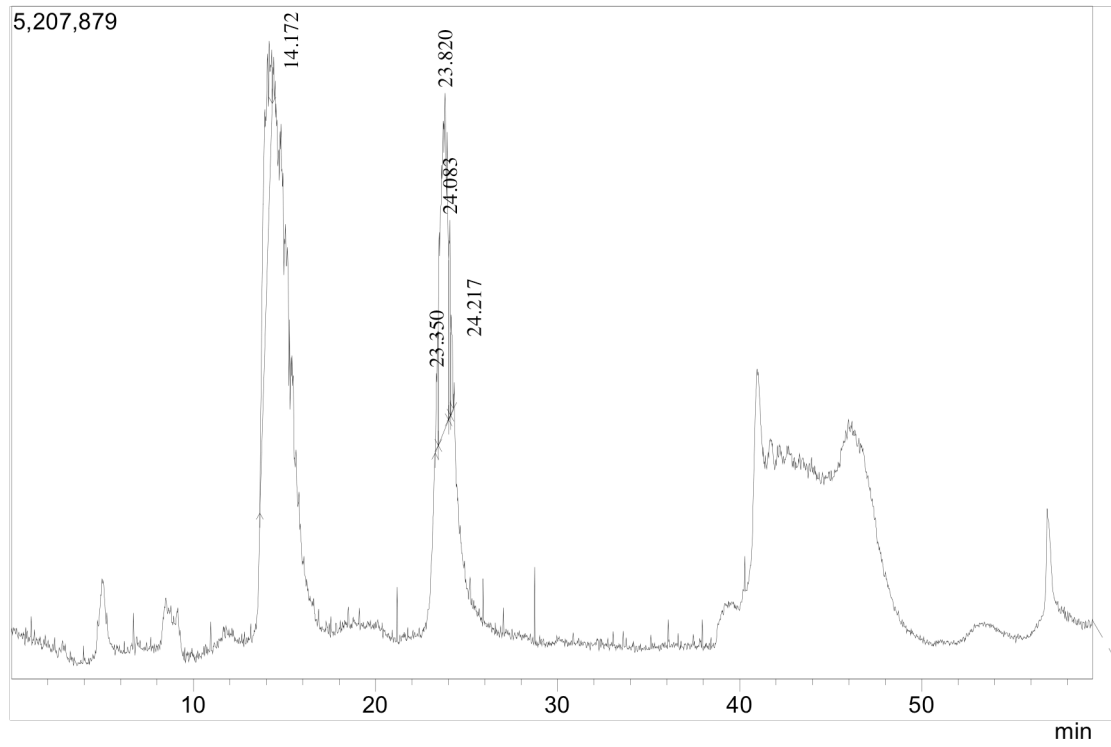


Figure A-15. Mass Data for pH 7.0 Aerobic Photolysis Run

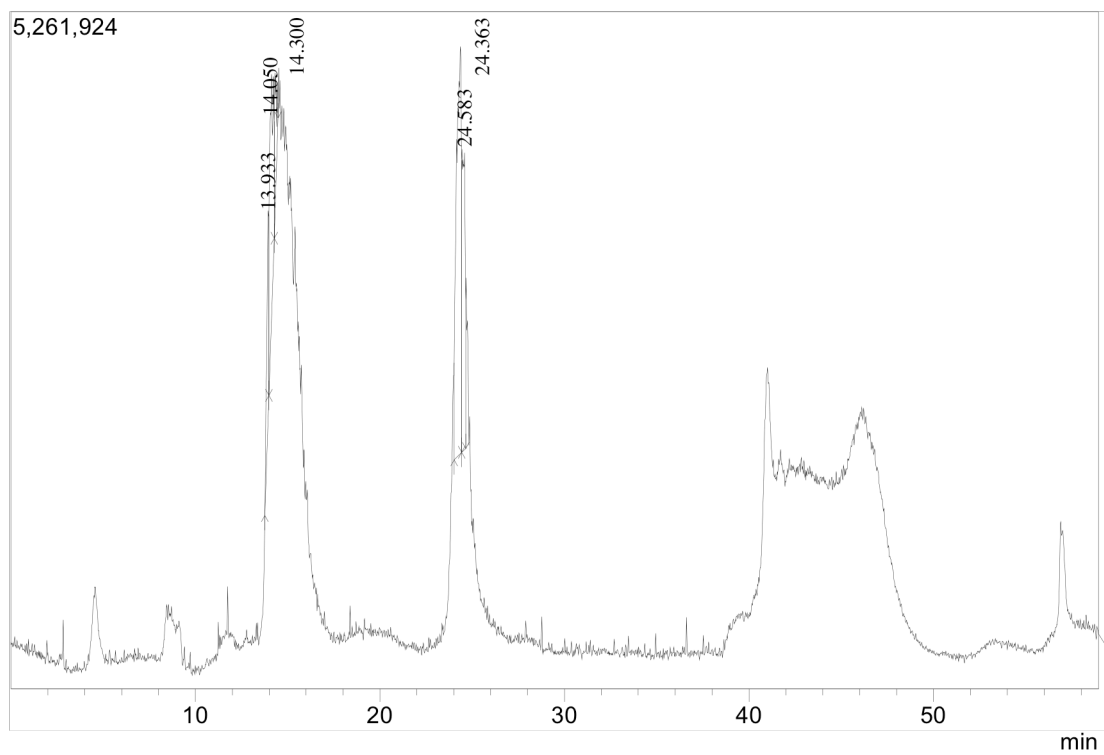
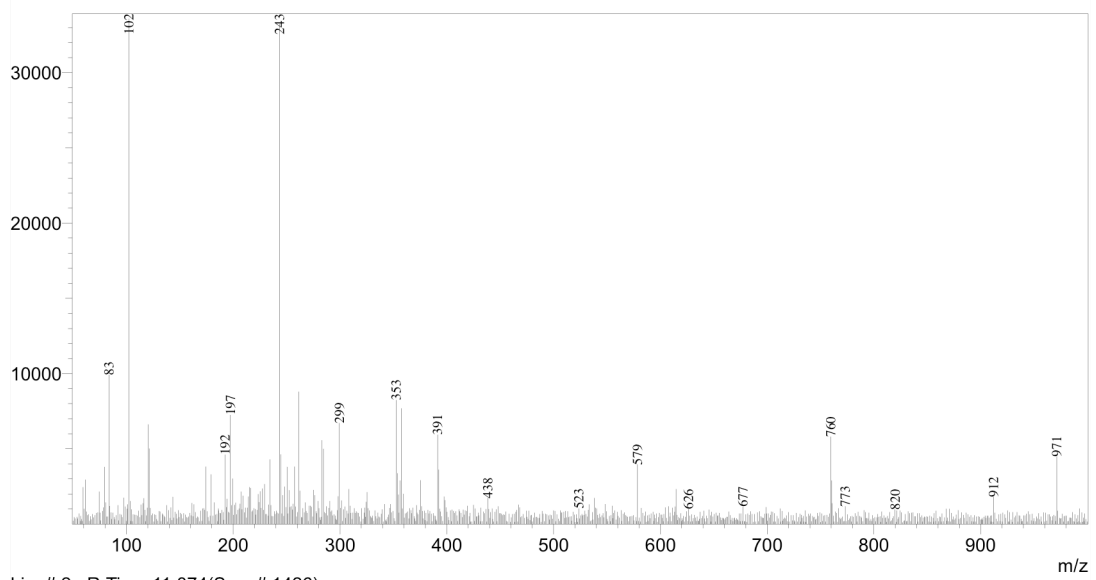


Figure A-16. Mass Data for pH 3.5 Aerobic Photolysis Run



Line#:2 R.Time:11.874(Scan#:1426)
 MassPeaks:1088
 RawMode:Single 11.874(1426) BasePeak:119(169583)
 BG Mode:None Segment 1 - Event 2

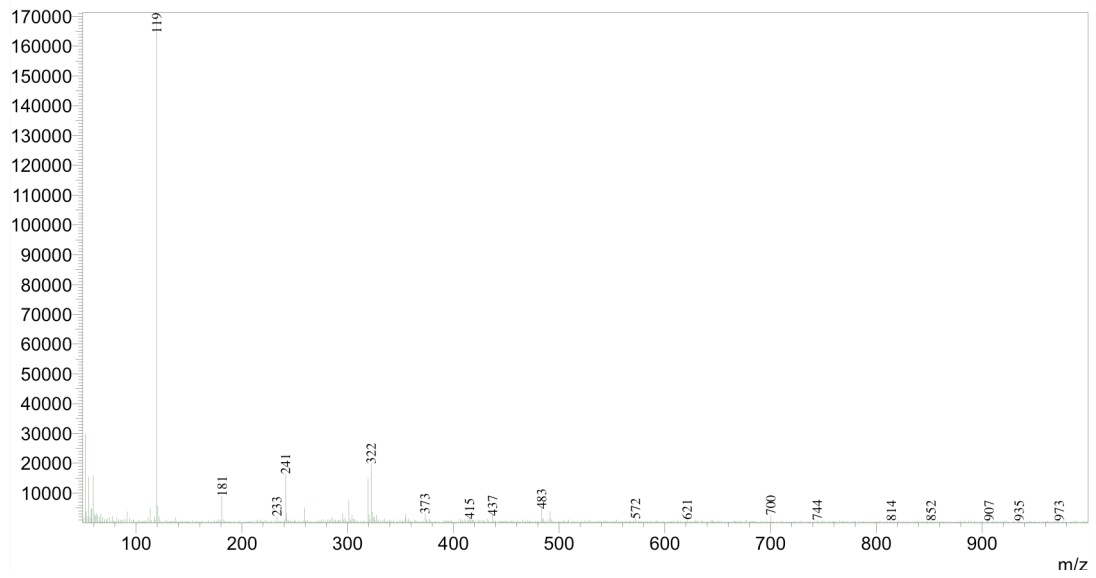


Figure A-17. Mass Data for 11.8 min, pH 7.0 Aerobic Photolysis Run

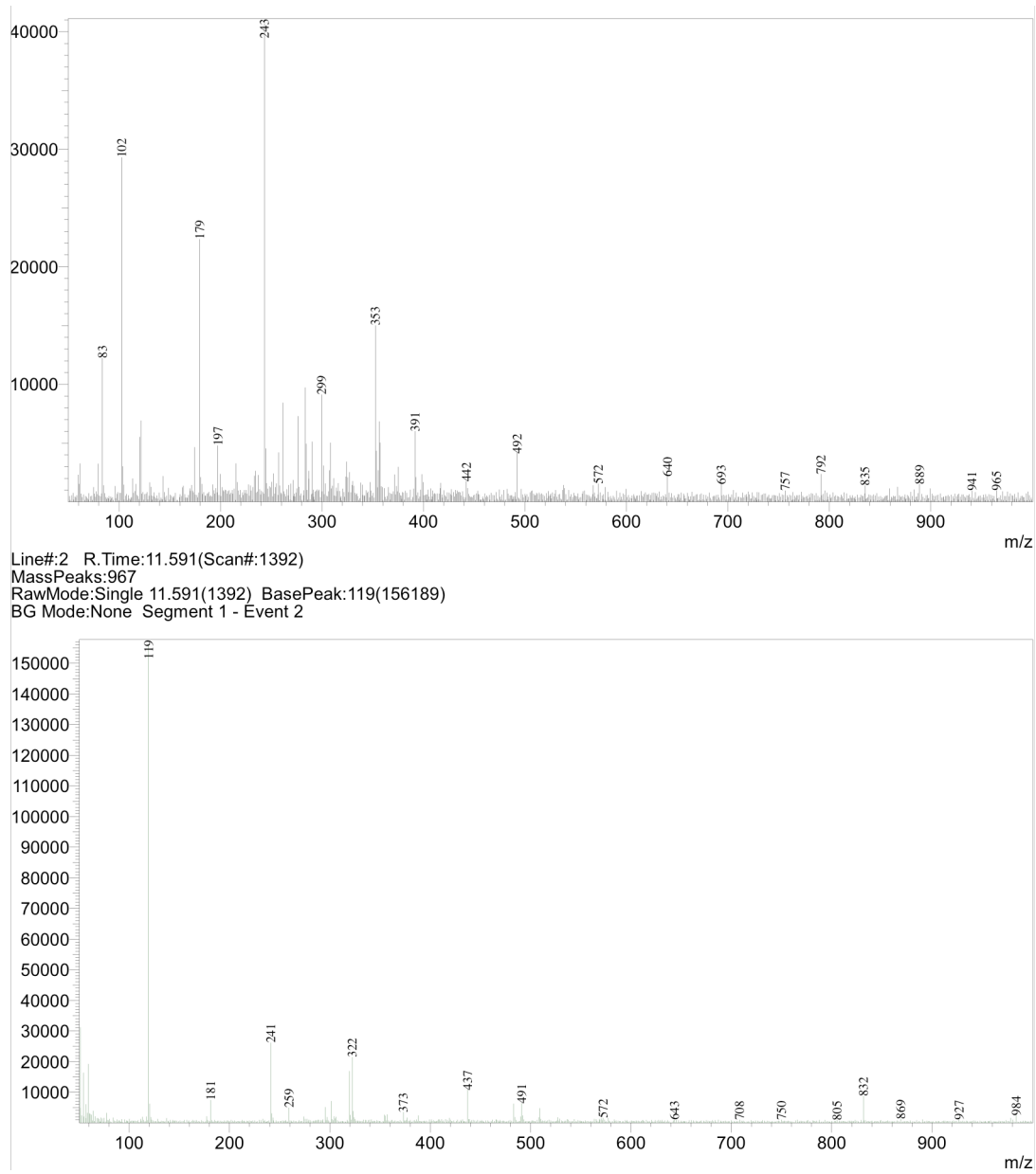


Figure A-18. Mass Data for 11.5 min, pH 3.5 Aerobic Photolysis Run

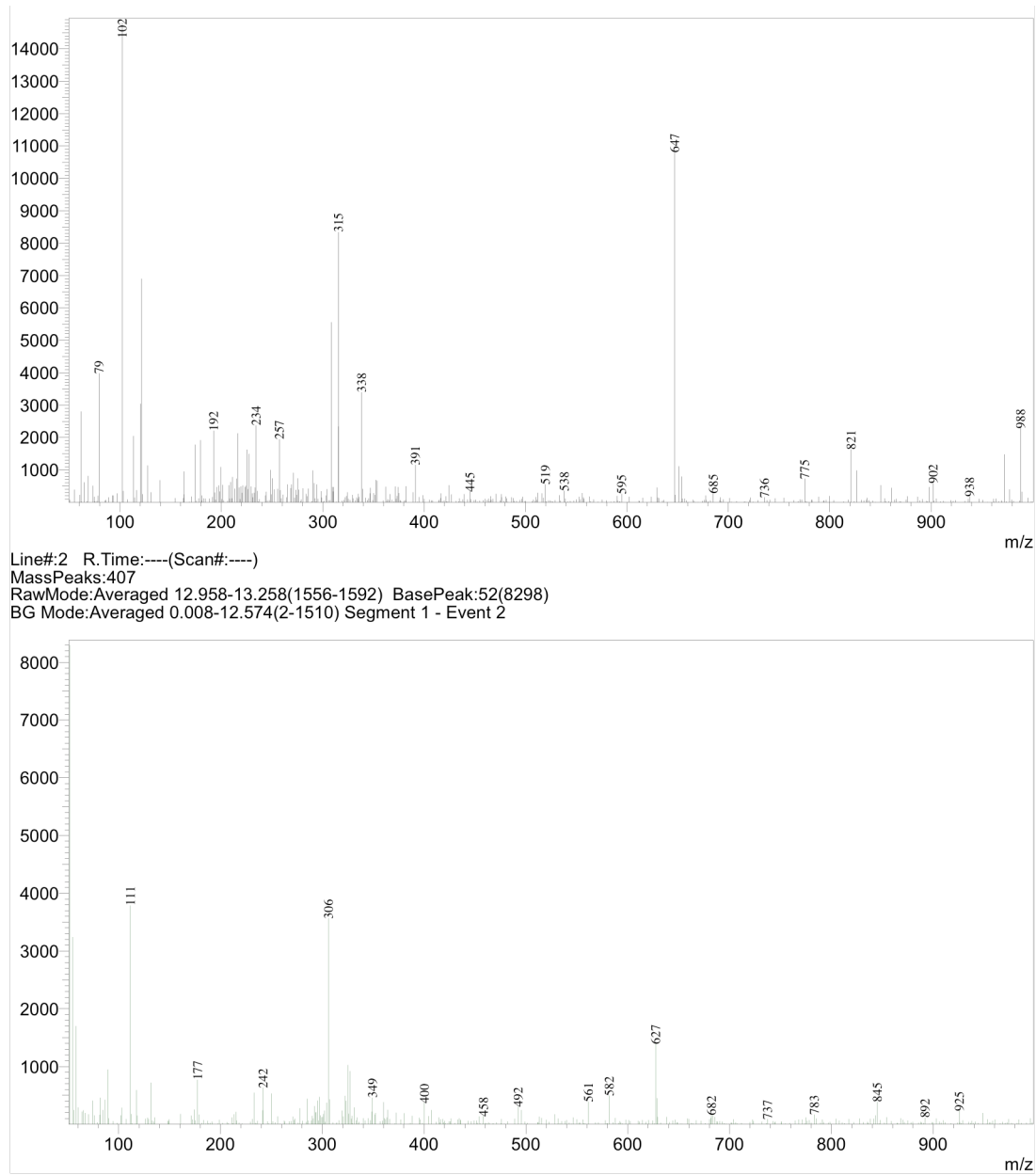


Figure A-19. Mass Data for 12.9-13.3 min, pH 7.0 Aerobic Photolysis Run

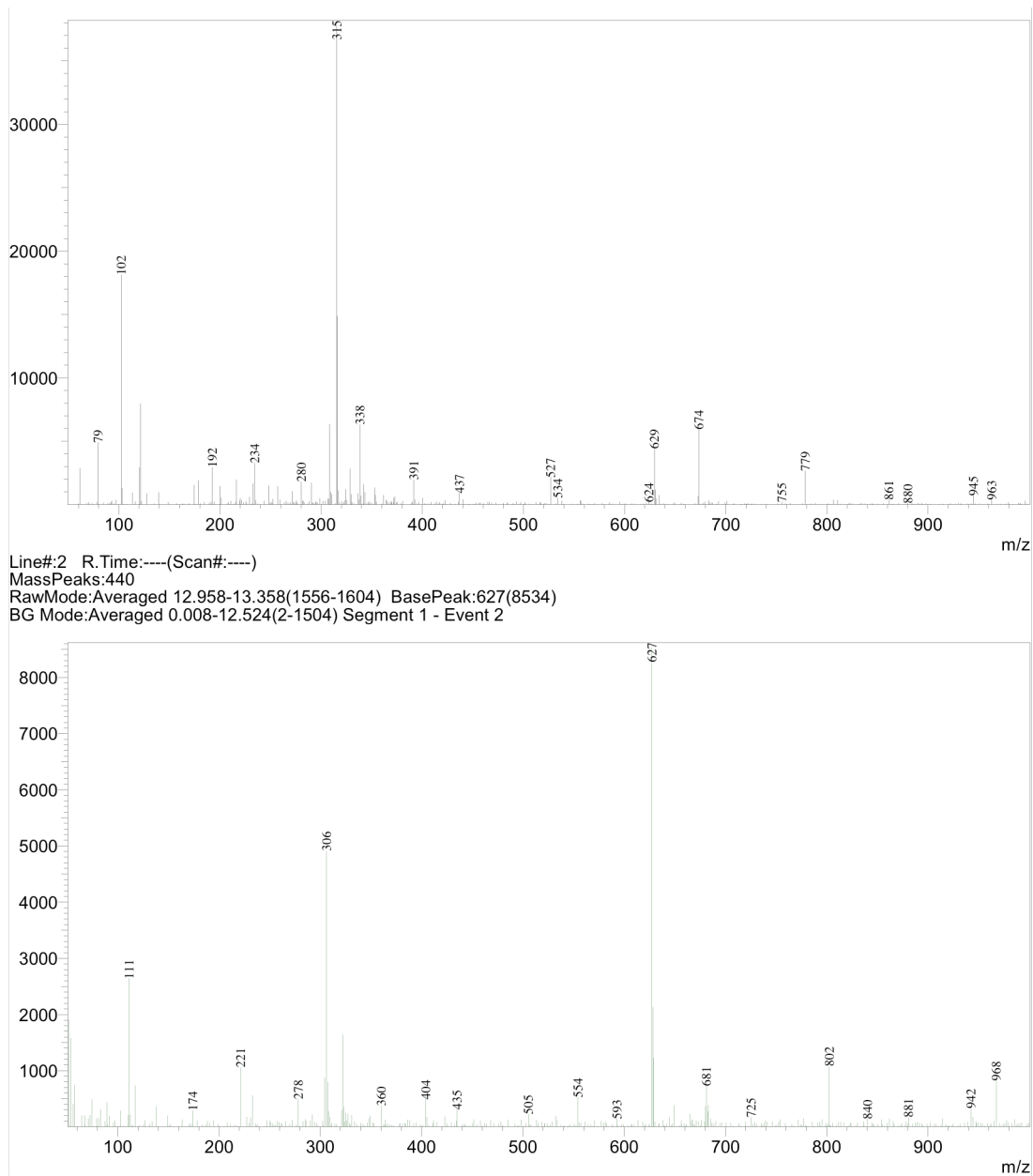


Figure A-20. Mass Data for 12.9-13.3 min, pH 3.5 Aerobic Photolysis Run

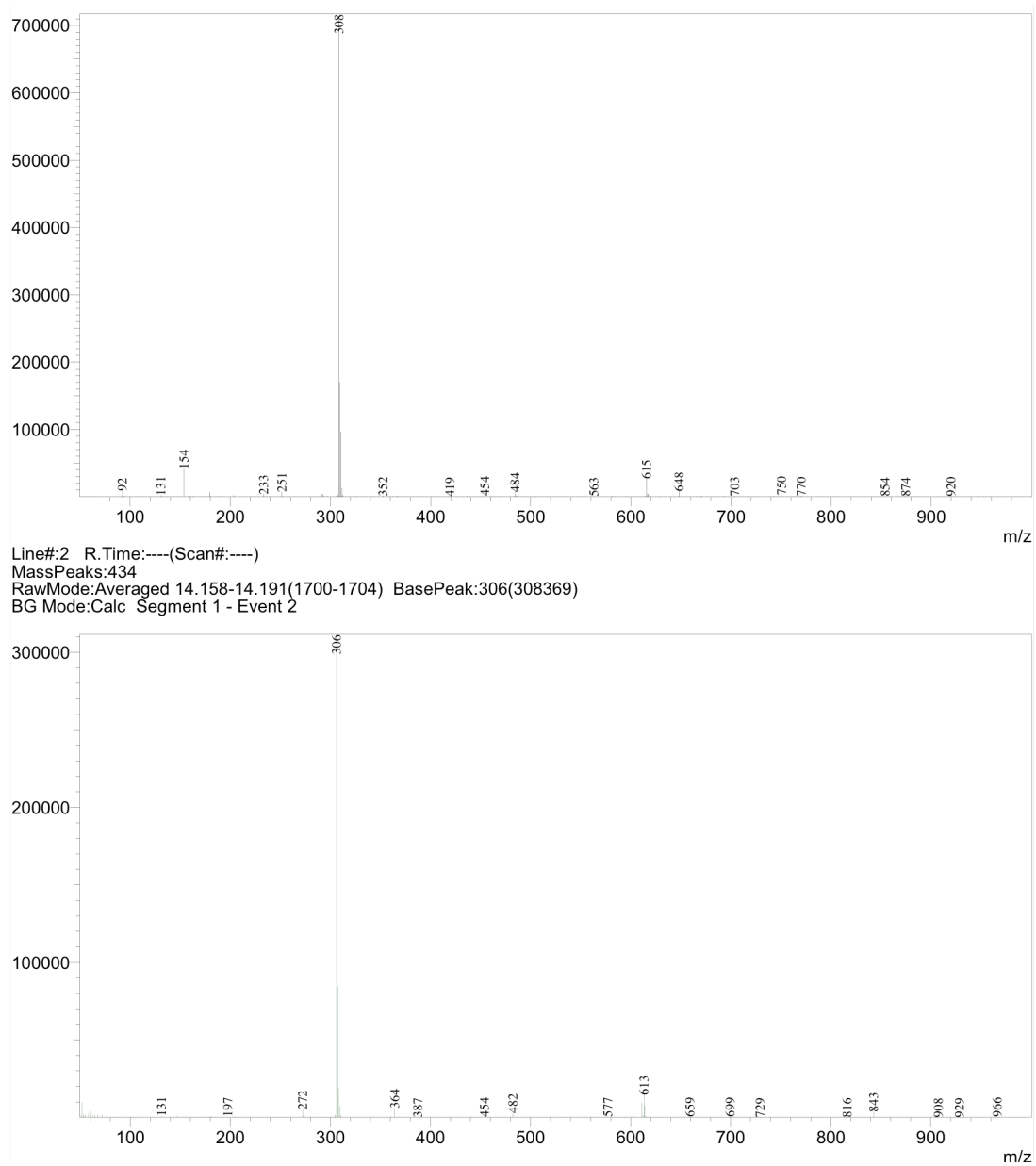


Figure A-21. Mass Data for 14.2 min, pH 7.0 Aerobic Photolysis Run

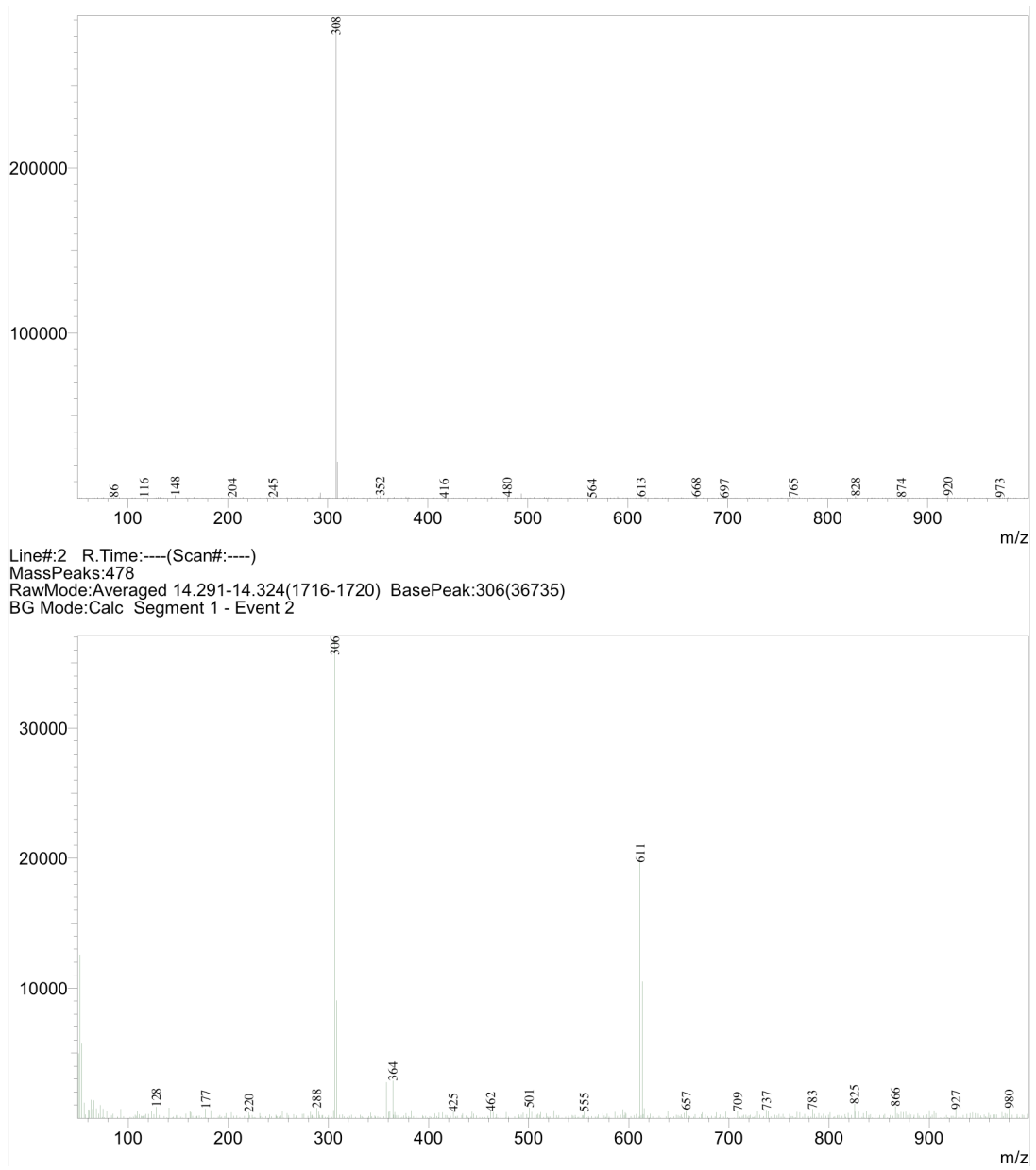


Figure A-22. Mass Data for 14.1 min, pH 3.5 Aerobic Photolysis Run

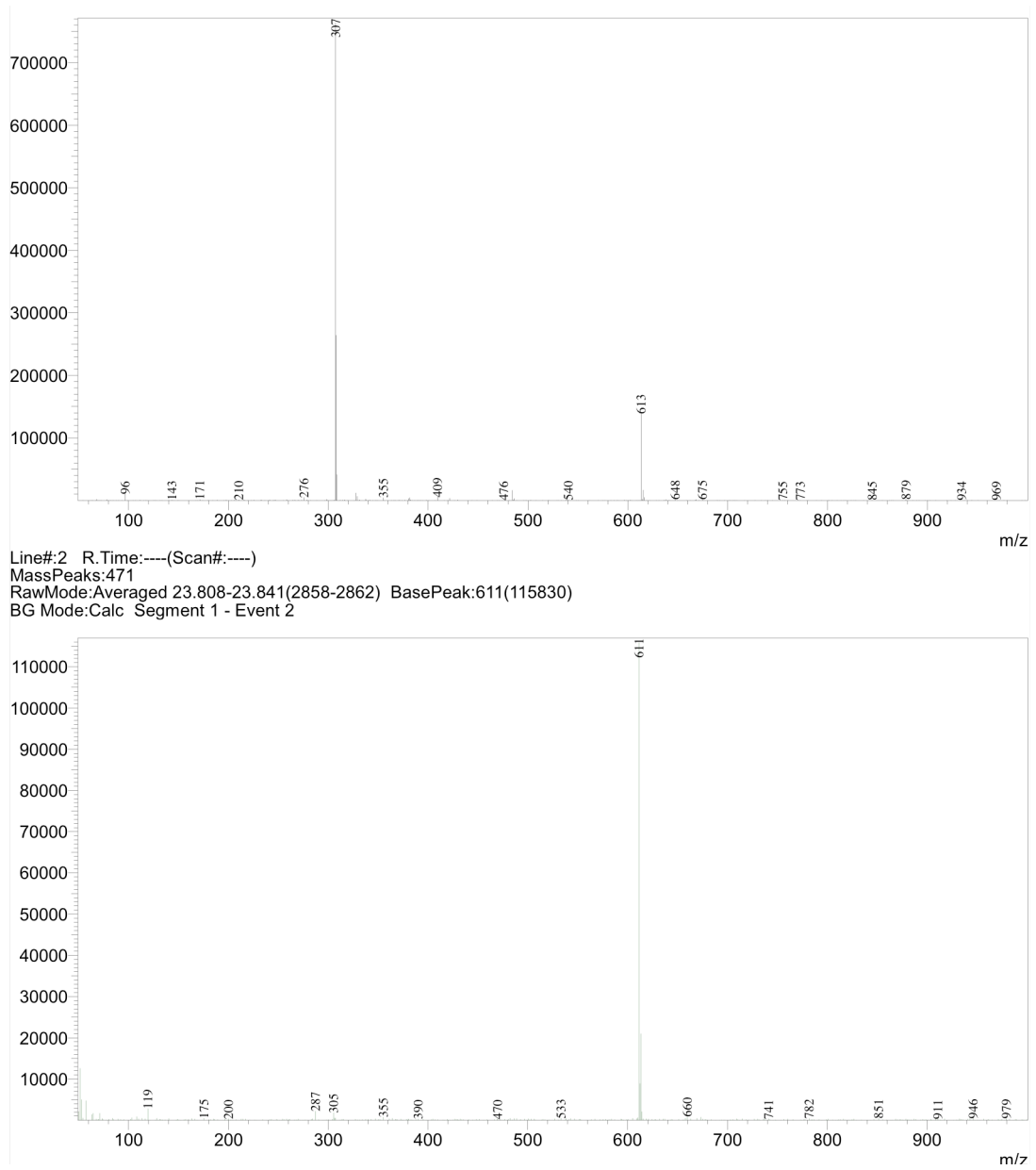


Figure A-23. Mass Data for 23 min, pH 7.0 Aerobic Photolysis Run

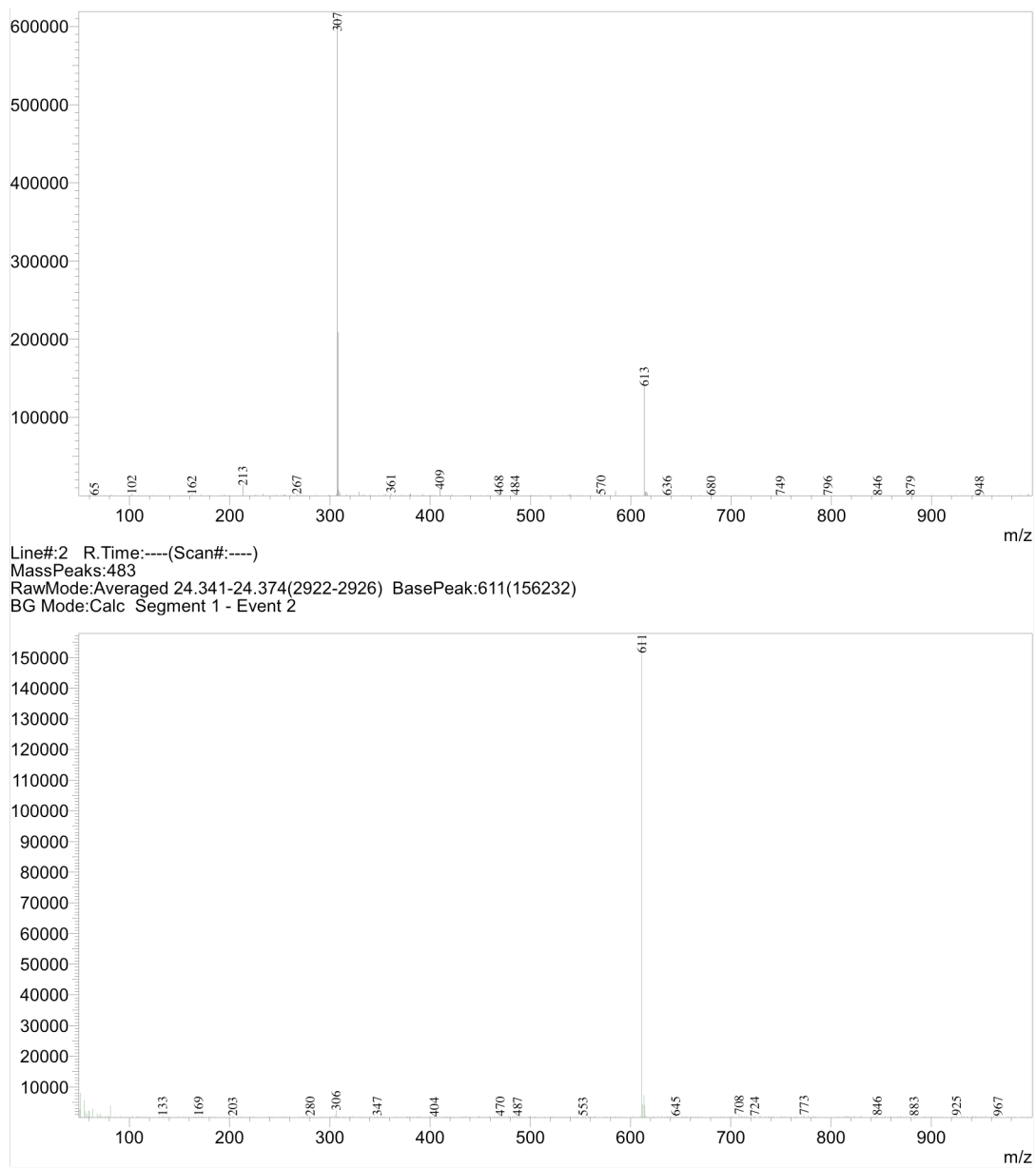
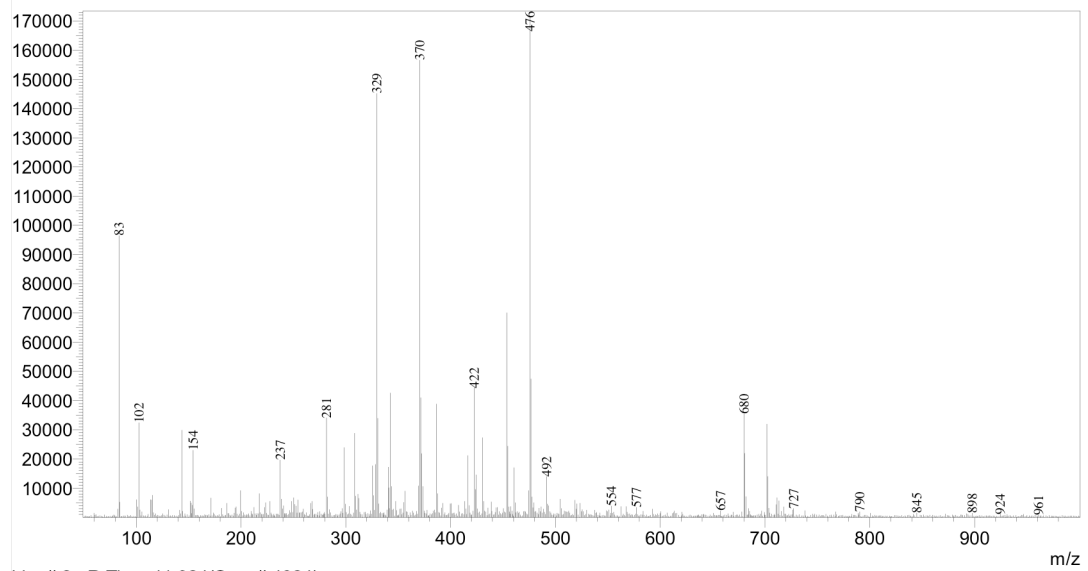


Figure A-24. Mass Data for 24 min, pH 3.5 Aerobic Photolysis Run



Line#:2 R.Time:41.024(Scan#:4924)
 MassPeaks:1003
 RawMode:Single 41.024(4924) BasePeak:387(541456)
 BG Mode:None Segment 1 - Event 2

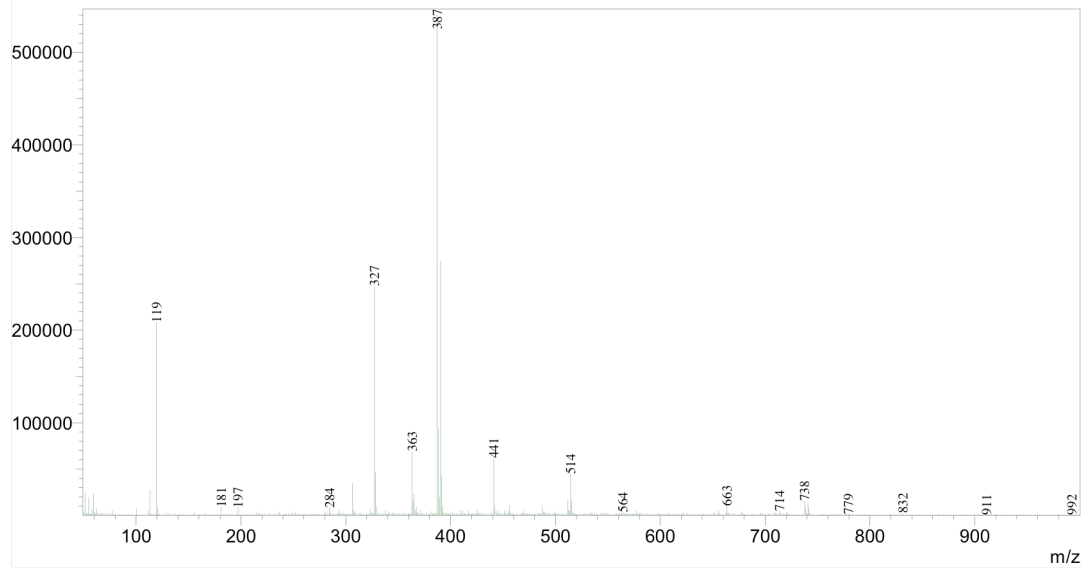
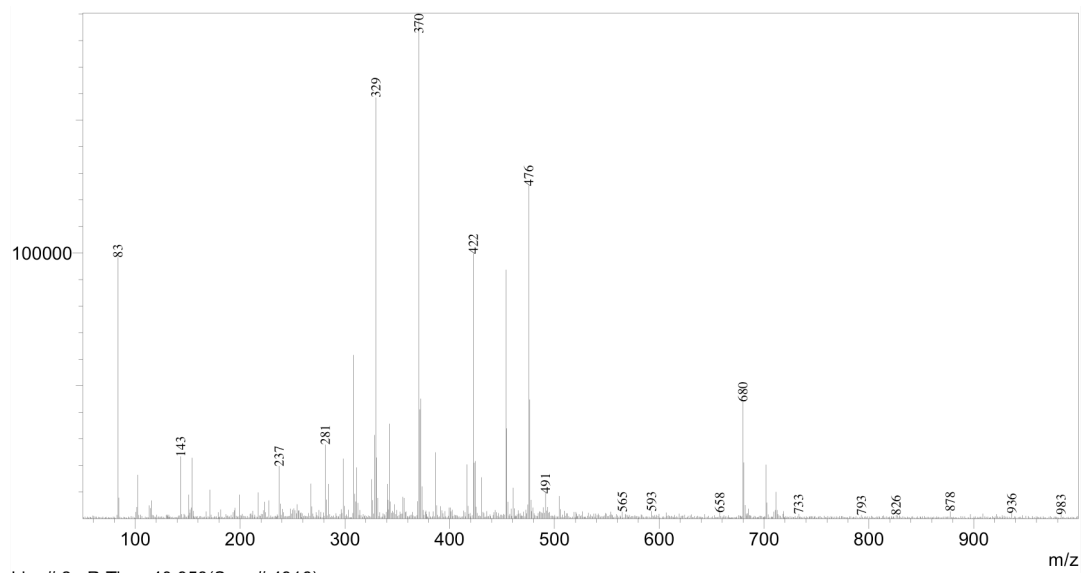


Figure A-25. Mass Data for 41.0 min, pH 7.0 Aerobic Photolysis Run



Line#:2 R.Time:40.958(Scan#:4916)
 MassPeaks:958
 RawMode:Single 40.958(4916) BasePeak:387(435518)
 BG Mode:None Segment 1 - Event 2

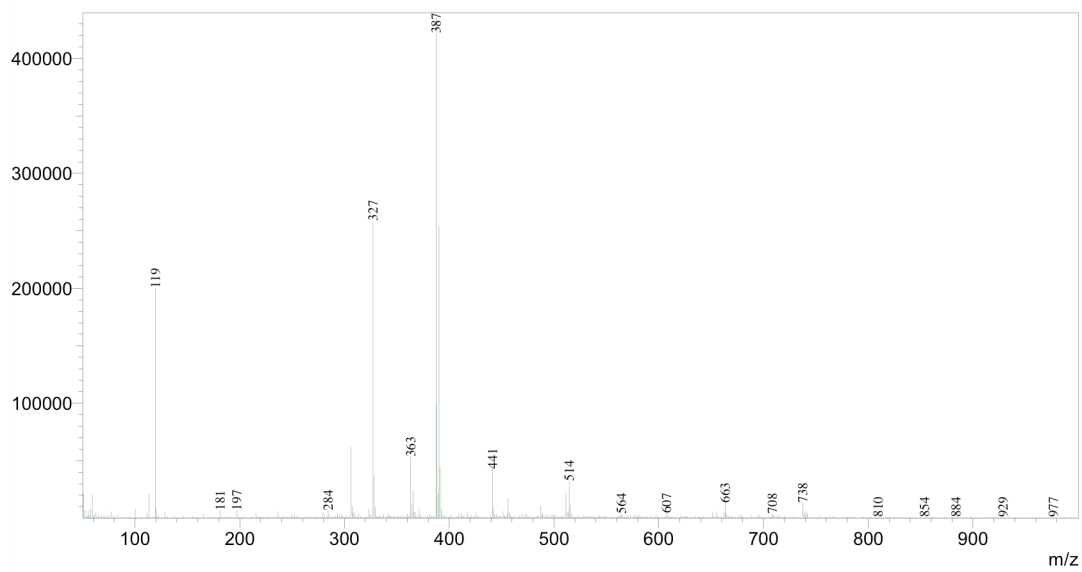


Figure A-26. Mass Data for 40.9 min, pH 3.5 Aerobic Photolysis Run

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